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SUMMARIES

RESULT 1 AX036745 LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE JOURNAL	C C 4-4-3 5-4-4-3	37 38 39 41 42	0000 0000 0000 0000	c c 22 c 22 c 25 24 29	10 11 11 11 11 11 11 11 11 11 11 11 11 1	© ® √ & W & W & P	Result No.
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synthetic construct.
synthetic construct
artificial sequences.
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Sequence 5
AX036739
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1 (bases 1 to 317)

Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from commelina yellow
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 3 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR);
GRUBER VERONIQUE (FR)
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/db_xref="taxon:32630"
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AX036738 348 bp
Sequence 4 from Patent W00058485.
AX036738
AX036738.1 GI:11226247
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Chimeric expression promotors originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 5 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIOUE (FR)
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l Similarity 98.4%;
62; Conservative
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                                                                                                                                                                                                                                                                1 (bases 1 to 398)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 6 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 (bases 1 to 348)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus patent: WO 0058485-A 4 05-CCT-2000;
MERISTEM HERAPBUTICS (FR); RANCE IANN (FR); THEISEN MANFRED; GRUBER YERONIQUE (FR)
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                                                                                                                                                                                                                                                                                                                                                                                          synthetic construct.
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340
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/db_xref="taxon:32630"
/note="promoter MPr1147"
                                                                                                                                                                                                                                                       Location/Qualifiers
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/db_xref="taxon:32630"
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80 с
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98.4%;
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Patent WO0058485.
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                                                                                                    Score 61.4; DB 6;
Pred. No. 5.6e-14;
0; Mismatches 1;
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Pred. No. 5.5e-14;
); Mismatches 1
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RESULT

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
Submitted (02-MAY-1990) Olszewski N.E., University of Minnesota,
Dept of Plant Biology, 220 Biological Sciences Center, 1445 Gort
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            artificial sequences.

1 (bases 1 to 243)

Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 1 05-027-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    243 bp
Sequence 1 from Patent WO0058485
AXO36735
                                            Direct Submission
Submitted (07-DEC-1990)
                                                                                                        Submitted (29-NOV-1990) Olszewski N.E., University of Minnesota, Dept of Plant Biology, 220 Biological Sciences Center, 1445 Gortner Ave., St. Paul, MN 55108, USA
                                                                                                                                                 3 (bases 1 to 7489)
Olszewski, N.E.
Direct Submission
                                                                                                                                                                                                                                           Medberry.S.L., Lockhart,B.E. and Olszewski,N.E. Properties of Commelina yellow mottle virus's complete sequence, genomic discontinuities and transcript sugges a pararetrovirus
                                                                                                                                                                                                                                                                                                                                                                                                                           complete genome; protease; reverse transcriptase; ril
Commelina yellow mottle virus
Commelina yellow mottle virus
Viruses; Retroid viruses; Caulimoviridae; Badnavirus
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                                                                           Olszewski, N.
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                                                                                                                                                                                                                                                                                                                    Ave., St. Pa
(revised by
                                                                                                                                                                                                                                                                                                                                    Dept of Plant Biology, 220 Bio
Ave., St. Paul, MN 55108, USA
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                               [3] The extra c at position 2672 permits readthrough to result in
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larity 100.0%;
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of Commelina Yellow Mottle
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/db_xref="taxon:32630"
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53 c
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s complete circular genome.
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suggest that
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163 kD ORF).
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GWGGEDNLLITRGLIGGRIANTSWOFAYDVRAWUELDSMGVKALKERKDAK
GWMLEPSKVVVPMOPTEMKAVSNVDGTTSLRESNYAAASTSKPPOYNEKDEEINKDEO
EINHSLLILLNDEESTDEDEEYYQYORX RAWGOVGDSTEYYDTDGVWEEIDRCDLPEY
VPSETSTPTIDESEAIIDEFLLHAYEORCDSDESLQSGDPRKYEYPTPOSSEHLDNE
SRSRSSSSTSTPTIDESEAIIDEFLLHAYEORCDSDESLQSGDPRKYEYPTPOSSEHLDNE
SRSRSSSASSTSMODDVEEIVRLMKEMRMKOOKKKAOQALSSOAQEEETIEEN EEN
KQAQEEPTQEEIPTHKENQPEEIQNEEILYFEEEPAFKHLAAQLDELVNMAESSGOSG
VGFOPPVNAQFDVNMEGPAGYAAPATSCATWSNCYNIPVKSANFRWKGPVGNFQLPSAQ
GKDGAMLVFGMNYSPEVFDRWASITRNYISSNNFNDGGDKIAMMEDLLGETERKIFVS
WRMRFKDEYQNLAKIANQDGGTQAILSOIRRIFIGADPVLGQNFVQNIFARKKCOLVC
PNYOSITRYLMDYLTLAAETCLMWSETEGPALSEELFTKMAPALGGRFVAOAYKLKDDT
SAVNLPSRVYFTINYLTEACTCLMWSETEGPALSEELFTKMAPALGGRFVAOAYKLKDDT
SAVNLPSRVYFTINYLTEACTCLMWSETEGPALSEELFTKMAPALGGRFVAOAYKLKDDT
SAVNLPSRVYFTINYLTEACTCLWSETEGPALSEELFTKMAPATGGLSROVG
PNFPGPAEVOMGVLKPKSSWRRPIQATLEEINCHNWTAISTGQLACRSCKQFLAGVQ
CHHCHAVYCFENCAEAYHDVQARKILSKDYSESARGKKGKAVIIEEDELEGEFLLSOLO
GENORLOKOVERLOEELMKLHREKDEAAKHSEKASRVFSTIQESDEAELNLIKEELRO
FKEETRMAIAQLKEAIIVQEEDTIEERCAMILEEKHTENIYSATAKAEYNGLYNWKVG
IKKDNMEPYYINAAIOTGATACLIOJSAIPERKENDAATACKAEYNGLYNWKVG
IKKDNMEPYYINAAIOTGATACLIOJSAIPERKENDAATACKAEYNGLYNWKVG
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                                                                     GRILIGEQYFRMPVTYVMNMGLSPGIQMIIGCSFIRSLEGGLRIEKDIITFYKLVTSI
ETSRTTQVANSIEELELSEDEYLNIAASVETPSFLDQEFARKNKDLLKEMKEMKYIGE
NPMEEWKNNKIKCKLNIINPDIKIMGRPIKHVTPGDEEAMTRQINLLLLQMKVIRPSES
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/translation="MSNITESGYKEALSVTNNYLAPAIGIGGATDVGLTGVTKOLNT
TIYLVAKITQOIEDLOSTIKRLESKGYKEALSVTNNYLAPAIGIGGATDVGLTGVTKOLNT
TIYLVAKITQOIEDLOSTIKRLESKGYKEALSVTNVTQDPNPEIISKLSDIQISLAR
QRAVNPAISGYSNYTAPTIKKVDRILRVFKKFN"
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1098..1505
/note="ORF 2"
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YSHRSEERMMVVNGSQVDRSFIOESSFEVLSRTGIEFIHIGVMLVRIQILHRKFAGTM
ALIVFRDTRWSDDRAVLAAMEIDLSEGNQIVYVLPDIMMTIKSFYRHIQICVMTKGYD
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/translation="MNVWLLKSHTPLGLLPYYSLLDPFCFMNQVDQVKQKLIDWLSSA
KLLSEEVYVFTPEVKINLRDLAHNIHIIAHRVALGFKVIYLVLVDIIFPLLKNIQKSQ
KELSEENLQSVLKIVKEQRRSLKQIEDQLSKVQSELAKLREDYLSRRPLSKQDVEELVV
RSKIYSKFDLKSGFWQVAMEEESVPWTAFLAGNKLYEWLVMPFGLKNAPAIFQRKMDN
                                KHRSTAFIVRSGTEIDPITGKEKKGKERMVFNYKLLNENTESDOYSLPGINTIISKVG
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/translation-"MATRRLPAVTQTDGSRTATESGVPEYEDQIRSYRNDQRRRHIWA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /protein_id="CAA37110.1"
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/note="ORF 3,
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/db_xref="GI:59049"
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/db_xref="GI:59048"
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/db_xref="taxon:10653"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         . 1098
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AX036755
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TITLE
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Matches 42; Conserv
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197 GGCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGTGGCATGCGTCATCCCTTACGT 256
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                           1 GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTT 60
                                                                                                                                                                                                                                                                                                                                                                                          Sequence 21
AX036755
AX036755.1
                                                         l Similarity
49; Conserv
                                                                                                                                                                                                                                                         artificial sequences.

1 (bases 1 to 392)
Rance, I., Theisen, M. and Gruber, V.
Rance, I. Theisen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 21 05-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                               synthetic construct.
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                                                                                                                                                        /organism="synthetic construct"
/db_xref="taxon:32630"
/note="promoter MPr1164"
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2977
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2672. .2674
/note="ccc was cc in [1]"
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CEAIIKFYNKTNENKPSRVRWLTFSDPLTGLGITVTFEHIDGKHNGLADALSRWINFI
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79.08;
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Pred. No. 9.4e
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; Pred. No. 6.6e-06;
0; Mismatches 0;
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AX036759
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                                                                                                                                                                                                1 (bases 1 to 472)
1 (bases 1 to 472)
1 (bases 1 to 472)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 25 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from commelina mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 23 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN M; GRUBER VERONIQUE (FR)
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Sequence 23 from Patent WO0058485.
AX036757
                                                                                                                                                                                                                                                                                                                                               Sequence 25 from Patent Ax036759
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                                                   Similarity
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                                       Conservative
                                                                                                  /note="promoter MPr1169"
1. .472
92 c 112 g 119
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/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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100.0%;
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95.2%;
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                                                   Score 37;
Pred. No.
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Pred. No. 9.5e-05;
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WO0058485
                                       Mismatches
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AX036758
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Rance, I., Thelsen, M. and Gruber, V.
Chimeric expression promoters originating importie virus and cassava vein mosaic virus Patent: WO 0058485-A 19 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (F); GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                             synthetic construct.
synthetic construct
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Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 24 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VEKONIQUE (FR)
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Ax036753
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1 (bases 1 to 541)
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/db_xref="taxon:32630"
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Search completed: May 24, 2003, 14:52:22 
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Sequence 20 from Patent WO0058485.
AX036754
AX036754.1 GI:11226263
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Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 22 05-CCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR); GRUBER VERONIQUE (FR)
LOCATION/Qualifiers
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Sequence 22 from Patent WO0058485.
AX036756
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Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 20 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR); GRUBER VERONIQUE (FR)
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35.2	35.2	35.2	35.2	35.2	35.2	35.2	35.2	35.2	35.2	35.2	35.2	35.2	35.2	35.6	35.6	35.6	35.9	36.2	36.2	36.8	36.8	37.1	38.1	39.0	39.0	39.4	39.4	39.4	39.4	. 39.4	56.2	56.2	56.2	58.7	58.7
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AAF59182	AAF59183	ABL90726	AAZ32842	AAX15104 ·	AAZ36238	AAA16693	AAF59199	AAL15716	ABK75564	AAH52365	ABA09445	AAH52940	AAA45410	AAK82827	ABN96872	ABL82625	ABL18414	AAC35439	AAC44959	ABL28884	AAH29763	ABN88295	ABL14808	AAC49145	AAC43051	AAN81551	AAS67507	ABK38141	AAF68230	ABL79935	AAA96856	AAA96854	AAA96853	AAA96858	AAA96859
gene	Human MyoD1 gene p	Human polynucleoti	Human high affinit	High affinity immu	cDNA encoding a bo	Human secreted pro	Human MyoDl gene p	Human breast cance	Bacillus lichenifo	S. epidermidis ope	Human IGERB homolo	S. epidermidis ope	Human secreted exp	Human immune/haema	Gene #3370 used to	Human ovarian canc	Drosophila melanog	Arabidopsis thalia	Arabidopsis thalia	Drosophila melanog	S cerevisiae apopt	Human large protei	Drosophila melanog			Bio F, Bio C and B	DNA encoding novel	cDNA encoding clon	Human lung tumour						Nucleotide sequenc

ALIGNMENTS

RESULT 1 AAA96845 Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; ss. Directional desoxynucleotide building block 19-FEB-2001 AAA96845; AAA96845 standard; DNA; 63 BP (first entry)

Synthetic.

WO200058485-A1.

05-OCT-2000

29-MAR-2000; 2000WO-IB00370

29-MAR-1999; 99FR-0003925

(MERI-) MERISTEM THERAPEUTICS.

WPI; 2000-647238/62. Rance I, Gruber V, Theisen

3

Chimeric expression promoter for transgenic plant production, comprises

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present sequence represents a directional desoxynucleotide building block, which was used to construct chimeric promoters of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably the first plant promoter originates from commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mossic virus. Especially, the promoters are
                                             Chimeric expression promoter for transgenic plant production, esequence from promoter comprising vascular expression region rewith sequence from promoter comprising green tissue expression
                                                                                                                                                                                                                                                                                      Chimeric -
                                                                                                                                                                                                                                                                                                                          Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression proplant green tissue expression promoter; Cassava vein mosaic
              Claim
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                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     promoter comprising vascular expression region replaced from promoter comprising green tissue expression region
                                                                                                                                                                           99FR-0003925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14 A; 14 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      23;
           91pp; English
                                                                                                                                                                                                                                                                                                                chimera;
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                                                                                                                        Theisen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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∠.8e-15;
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mosaic virus;
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Best Local S
Matches 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergent regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                 Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric - Commelia
Chimeric - Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus transgenic plant; chimera; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA96839 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200058485-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTT
                                                                                                                                                                                                                                                                                                                                                                                                               2000-647238/62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCCAAGCTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Commelina yellow mottle virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B₽;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000WO-IB00370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99FR-0003925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        of chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The1sen
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 74 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 75 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus;
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The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from

The present sequence represents a chimeric promoter of the invention The specification describes chimeric expression promoters. These

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Best Local :
                                                                                                      The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Chimeric - Commelina yellow mottle virus Chimeric - Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Commelina yellow mottle virus, and the from the Cassava vein mosaic virus. Fearived from intergenic regions. The
                                                                                                                                                                                                                                                                                                                                                                                                    Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric expression promoter; plant green tissue expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAA96841 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-647238/62.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rance
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (MERI-) MERISTEM THERAPEUTICS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          309
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            61 TAT 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCCGGTTCCCAAGCTT 60
                                                                                                                                                                                                                                                                                                              present sequence represents a chimeric promoter of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             eric expression promoter for transgenic plant production, comprises ence from promoter comprising vascular expression region replaced sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TAT 311
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTT 308
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                                                                                        from
                                                                                                                                                                                                                                                                                                                                                                                                    Page 82;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                       intergenic regions.

y transgenic plants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        99FR-0003925
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                              91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%;
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Pred. No. 4e-
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plant vascular expression promoter; promoter; Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              89 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     expression promoter
                                                                                  The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4e-15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MPr1154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Sequence

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RESULT 5
                                                                     Query Match
Best Local S
Matches 62
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Best Local
                                                                                                                                                                                      The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant green tissue expression commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chimeric sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric expression promoter; plant vascular expression plant green tissue expression promoter; Cassava vein motransgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chimeric - Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA96838
                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-647238/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-OCT-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200058485-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA96838;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MERI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        61
                                                                                                                                                                        producing
eric expression promoter for transgenic plant production, cence from promoter comprising vascular expression region resource from promoter comprising green tissue expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAT 243
                                                                                                                                                                                                                                                                                                                                                                                <u>ن</u>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ı,
                                                                     62; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MERISTEM THERAPEUTICS.
                                                                                                                                         348
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          63
                                                                                                                                                                                                                                                                                                                                                                                Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Commelina yellow mottle virus Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sequence of chimeric expression promoter MPr1117
                                                                     Conservative
                                                                                                                                         BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000WO-IB00370
                                                                                                                                                                        transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                              81; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         99FR-0003925.
                                                                                                                                      116 A; 70 C; 78 G; 84 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  entry)
                                                                                    97.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97.5%;
98.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Theisen M;
                                                                     0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 61.4; D
Pred. No. 1.6e
0; Mismatches
                                                                   Score 61.4; D
Pred. No. 1.7e
0; Mismatches
                                                                                    61.4; DB 21
No. 1.7e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1.6e-14
                                                                                                                                      0 other;
                                                                                                    21;
                                                                     1;
                                                                     Indels
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                                                                                                     Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  region replaced
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                301;
                                                                                                       348;
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                                                                                                                                                                                                                                                                                                                                                                                                                                  region
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RESULT 6
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                                                                                                                            Query Match
Best Local :
                                                                                                              Matches
                                                                                                                                                                                                     The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                    Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric -
Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA96840 standard; DNA; 398
                                                                                                                                                                   Sequence 398
                                                                                                                                                                                                                                                                                                                                                                 Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200058485-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA96840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (MERI-) MERISTEM THERAPEUTICS
 338
                                                       278
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           288
                           61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
                                                                                                                           Local
                                                                                  H
                                                                                                                                                                                             producing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAT
TAT 340
                           TAT 63
                                                                    GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TAT 290
                                                       GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCTGGTTCCCCAAGCTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                              2000-647238/62
                                                                                                                                                                                                                                                                                                                                                                5; Page 82;
                                                                                                                           Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Commelina yellow mottle virus. Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      sequence of
                                                                                                                                                                BP; 128 A; 80 C; 93 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2000WO-IB00370
                                                                                                                                                                                            transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99FR-0003925
                                                                                                                                                                                                                                                                                                                                                                91pp; English
                                                                                                                        97.5%;
98.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Theisen M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    chimeric expression promoter MPr1147.
                                                                                                           ; Score 61.4; D
; Pred. No. 1.7e
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ВP
                                                                                                                                                                97 T; 0 other;
                                                                                                                         DB 21;
L.7e-14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               vein mosaic virus;
                                                                                                            Indels
                                                                                                                                      Length
                                                                                                           0;
                                                                                                           Gaps
                                                                                                            0
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RESULT

chimeric Promoter;

expression

intergenic region; Commelina yellow mottle virus; expression promoter; plant vascular expression pro

promoter;

Nucleotide sequence of chimeric expression promoter MPr1164.

19-FEB-2001 AAA96855;

(first

entry)

AAA96855 standard;

DNA;

BP.

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RESULT 8
AAA96855
ID AAA9
XX
AC AAA9
XT 19-F
XX
DE Nucl
XX
KW Prom
KW Chim
                                                                                                                                                                                         В
                                                                                                                                                                                                                                                        Query Match
Best Local S
                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents a promoter fragment from the intergenic region of commelina yellow mottle virus. The promoter is a strong promoter in vascular and reproductive tissues. The promoter is used to construct chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            plant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                     promoter comprising a plant green tissue expression promoter reg
Preferably, the first plant promoter originates from Commelina y
mottle virus, and the second plant promoter originates from the
vein mosaic virus. The chimeric promoters are useful for produci
                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                            transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAR-2000; 2000WO-IB00370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200058485-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Commelina yellow mottle virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric expression promoter; plant green tissue expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA96835;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA96835 standard; DNA; 243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                         153
                                                                                                                                                                                                        22 TTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTTTAT 63
                                                                                                                                                                                          TTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTTTAT 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          H,
                                                                                                                                                                                                                                          l Similarity
42; Conserv
                                                                                                                                                                                                                                                                                                 243 BP; 71 A; 53 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gruber V,
                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    79;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 intergenic region of Commelina yellow
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99FR-0003925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91pp;
                                                                                                                                                                                                                                                        66.7%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Theisen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    English.
                                                                                                                                                                                                                                             0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ₽P
                                                                                                                                                                                                                                                                                                 45 G;
                                                                                                                                                                                                                                                         Score 42;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           plant vascular expression
                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                 74 T; 0 other;
                                                                                                                                                                                                                                                         DB 21; 1
7.2e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cassava vein mosaic
                                                                                                                                                                                                                                             0,
                                                                                                                                                                                                                                                                      Length 243;
                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mottle virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           promoter;
                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                                                                                                                                                 region.
                                                                                                                                                                                                                                                                                                                                                      yellow
e Cassava
                                                                                                                                                                                                                                              Gaps
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ARASSILT 9
ARASSICT 10
ARASSICT ARASSIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
                                                                Chimeric - 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Chimeric sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoters are derived from intergenic regions. The chimeric promoters are useful
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                                                                                                                                                                                                  chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chimeric -
Chimeric -
   WO200058485-A1
                                                                                                                                                                         transgenic
                                                                                                                                                                                                                                                                             Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                              Nucleotide sequence of chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                      19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA96857;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAA96857 standard; DNA; 604 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 392 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (MERI-)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200058485-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           plant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              eric expression promoter for transgenic plant production, comprisence from promoter comprising vascular expression region replaced sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GCCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTT 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2000-647238/62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Page 86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gruber V,
                                                                Commelina yellow mottle virus Cassava vein mosaic virus.
                                                                                                                                                                         plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Commelina yellow mottle virus Cassava vein mosaic virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000WO-IB00370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                127 A; 80 C; 87 G; 98 T; 0 other;
                                                                                                                                                                     chimera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    91pp; English.
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                                                                                                                                                                         88
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Pred. No. 1.6e
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                           promoter MPr1167
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RESULT 10
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Best Local Similarity
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                                                                                                                                                                                                                                                                    Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA96859 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim
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Rance I,
                                                                                 29-MAR-1999;
                                                                                                                       29-MAR-2000;
                                                                                                                                                                05-OCT-2000.
                                                                                                                                                                                                         WO200058485-A1.
                                                                                                                                                                                                                                                 Chimeric - Cassava
                                                                                                                                                                                                                                                                                                         chimeric expression promoter; plant plant green tissue expression promot transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                         Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                Nucleotide sequence of chimeric expression promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA96859;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  05-OCT-2000
                                        (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   for producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              278 GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCTAGT 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     \vdash
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      eric expression promoter for transgenic plant production, ence from promoter comprising vascular expression region requence from promoter comprising green tissue expression
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Gruber V,
                                                                                                                                                                                                                                                                    Commelina yellow mottle virus
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                                                                                                                       2000WO-IB00370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA;
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95.2%;
Theisen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            472
                                                                                                                                                                                                                                                 mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
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Pred.
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No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    genic plant production, comprises ular expression region replaced green tissue expression region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21;
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                                                                                                                                                                                                                                                                                                                                                                                                                MPr1169.
                                                                                                                                                                                                                                                                                                                                mosaic virus;
                                                                                                                                                                                                                                                                                                                                                      promoter;
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RESULT 11
RABAS6858
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Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                        with sequ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
The present sequence represents a chimeric promoter of the The specification describes chimeric expression promoters.
                                                                                Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chimeric -
Chimeric -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transgenic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAA96858;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAA96858 standard; DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 5;
                                                                                                                                                                                                                                                                                                                                                                (MERI-) MERISTEM THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                          Page
                                                                                                                                                     expression promoter for transgenic plant production, comprises from promoter comprising vascular expression region replaced uence from promoter comprising green tissue expression region
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from promoter comprising vascular expression region replaced
uence from promoter comprising green tissue expression region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    472 BP; 149
                                                                                                                                                                                                                                                                                                             Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence of chimeric expression promoter MPr1168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Commelina
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                                                                                                                                                                                                                                                                                                                                                                                                                   99FR-0003925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chimera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          91pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      vein mosaic virus
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                                                                        91pp; English.
                                                                                                                                                                                                                                                                                                             Theisen M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 yellow mottle virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; 92 C; 112 G; 119 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    541
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plant vascular expression promoter;
promoter; Cassava vein mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Pred.
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  invention These
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RESULT 12
AAA96853
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Best Local :
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                                       The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originate from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                     Chimeric expression promoter for transgenic plant production, compris sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                           Claim
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Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric expression promoter; plant green tissue expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleotide sequence of chimeric expression promoter MPr1162
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200058485-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             for producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                278 GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local
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                               producing
                                                                                                                                                                                                                        5; Page 85; 91pp; English
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37; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Commelina yellow mottle virus Cassava vein mosaic virus.
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                             transgenic
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       393
                             plants.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0;
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 21;
. 8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    314
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saic virus;
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Sequence

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RESULT 13
AAA96854
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                                                   Matches
                                                                Query Match
Best Local
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                                                                                                                                  The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                    Chimeric expression promoter for transgenic plant production, compris sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
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Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus;
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                                                                                                   Sequence 462
                                                                                                                                                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                             WPI;
                                                                                                                                                                                                                                                                                                                                                                                                     Rance
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleotide sequence
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                                                                                                                            for producing
                                                                                                                                                                                                                                                        The present sequence represents a chimeric promoter of the invention
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  197
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              GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC
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GCCGGAAAGTAACCTTATGCATTTGTAATTTGGTTAC
                                                                                                                                                                                                                                                                                                                                                                             2000-647238/62
                                                                                                                                                                                                                                                                                   5; Page 86;
                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                               MERISTEM THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Commelina yellow mottle virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               plant; chimera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                   BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first
                                                                                                                            transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                       99FR-0003925
                                                                                                     148
                                                                                                                                                                                                                                                                                   91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           vein
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                                                                                                  A; 87
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97.3%;
                                                            56.2%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           mosaic virus.
                                                                                                                            plants.
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                                               Score 35.4; E
Pred. No. 0.00
0; Mismatches
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Pred. No. 0.00
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                    X
                                                                                                  111 G; 116 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; DB 21;
0.00032;
ches 1;
                                                            ,00033;
                                                                        DB 21;
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                                                                                                                                                                                                                                                                                                                        tissue expression region
                                                                        Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length
                                                                          462;
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RESULT 15
ABL79935/c
ID ABL799
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AC ABL799
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AC ABL799
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AC T7-MAY
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DT 17-MAY
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KW Human;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimeric expression promoter; plant vasue plant green tissue expression promoter; transgenic plant; chimera; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric - Commelina yellow mottle virus. Chimeric - Cassava vein mosaic virus.
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                                                                                                                         ABL79935
                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                               derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence
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                                                                                                                                                                                                                                                               Local
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                                                                                                                                                                                                                                                                                                                                  producing
                                                                                                                                                                                           GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          eric expression promoter for transgenic plant production, cence from promoter comprising vascular expression region researce from promoter comprising green tissue expression
                                                                                                                                                                                                                                                 36; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 86-87; 91pp; English
                                                                                                                         standard;
                                                                                                                                                                                                                                                                                                        600 BP;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence of chimeric expression
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                                                                                                                                                                                                                                                                                                                                  transgenic plants.
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                                                                                                                                                                                                                                                                                                      188 A; 111 C; 147 G; 154 T; .0 other;
                                                                                                                         CDNA; 434
                                                                                                                                                                                                                                                           56.2%;
97.3%;
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Pred. No. 0.00
0; Mismatches
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Human ovarian cancer related cDNA clone SEQ ID NO:2913.

(first entry)

Human; ovarian cancer; ovarian tumour;

cytostatic; gene;

SS

Homo sapiens.

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                                                                                                                                                                             The present invention describes a composition (I) comprising: carriers CC and immunostimulants; and a polypeptide (II) of a ovarian tumour CC polypeptide encoded by a polynucleotide (II) having a cDNA sequence CC (S1) from the 10912 nucleotide sequences as given in ABL77023 to CC ABL87934, (III) encoding (II) having a sequence (S2), a T cell CC population of (II), or antigen presenting cells that express (II). CC (I) has cytostatic activity. An oligonucleotide (IV) that hybridises to CC (S1) can be used for detecting ovarian cancer in a patient's biological CC sample preferably serum or ovarian tissue. The method comprises CC contacting a biological sample from a patient with (IV), detecting the CC amount of polynucleotide hybridising to (IV) and comparing the amount to polynucleotide hybridising to (IV) and comparing the amount to patient, where the amount of polynucleotide hybridising to (IV) is Cdetected preferably by polymerase chain reaction (PCR). (I) comprising CC (III) and/or (II) is useful for stimulating and/or expanding T cells specific for an ovarian tumour protein comprising contacting T cells cand protein or (II). (III) is useful in design and preparation of cand proteins in tumour cells; and to isolate a full length gene from a contacting tell in the contacting in tumour cells; and to isolate a full length gene from a contacting e.g., a tumour cDNA library using well known contacting tells.
                                                                                     Query Match
Best Local S
Matches 38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID 2913; 489pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Composition for therapy and diagnosis of ovarian cancer comprising polypeptide of a ovarian tumor polypeptide, polynucleotide encoding polypeptide, antibody specific to polypeptide or T cell expressing polypeptide -
                                                                                                                                                                      Sequence 434 BP; 136 A; 66 C; 92 G; 140 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-122075/16.
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                                                                                                                                                                                                                   techniques.
92 GGTAAGCAACATTGTTCATTTGTTACTGGGTGTTCTAGATCAAACCTTCACAAGCTATAT 33
                        4 GGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCCAAGCTTTAT 63
                                                                                     l Similarity
38; Conservat
                                                                                       Conservative
                                                                                   39.4%; Score 24.8; I
63.3%; Pred. No. 5;
tive 0; Mismatches
                                                                                                                           DB 24; Length 434;
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Search completed: May 24, 2003, 14:34:43 Job time : 110.726 secs

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Post-processing: Minimum Match 0%
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Listing first 45 summaries
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1: /cgn2_6/ptodata/1,
2: /cgn2_6/ptodata/1,
3: /cgn2_6/ptodata/1,
4: /cgn2_6/ptodata/1,
5: /cgn2_6/ptodata/1,
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 ggcggaaagtaaccttatgc.....tgccggttcccaagctttat
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                                                                                                                                                                                                                                                                                                                                                                     /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
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US-08-727-118-1
US-08-680-506-4
US-08-680-506-2
US-08-680-506-1
US-08-680-506-1
US-08-680-506-2
US-08-444-803-36
US-08-445-265A-36
US-08-445-265A-36
US-08-445-265A-36
US-08-455-244-36
US-08-455-244-36
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US-08-455-265A-36
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US-08-916-902A-2
US-09-213-389-2
US-09-193-792-2
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        8-181-271A-36
8-449-315-36
8-444-803-36
8-449-043-36
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                                                                                                                Sequence 2, Application Patent No. 5871930 GENERAL INFORMATION:
                                                                                                                                                                                                                              Query Match 35.3
Best Local Similarity 64.7
Matches 33; Conservative
                                                       APPLICANT: Bandman, Olga
APPLICANT: Lal, preeti
APPLICANT: Corley, Neil C.
TITLE OF INVENTION: HIGH AFI
TITLE OF INVENTION: RECEPTO)
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
                                                                                                                                                                                        LENGTH: 1312
                                             CORRESPONDENCE ADDRESS
       CITY: Palo Alto
STATE: CA
                            STREET:
COUNTRY:
                                      ADDRESSEE:
                                                                                                                                       Application US/08916902A
                             3174 Porter Drive
USA
                                      Incyte Pharmaceuticals, Inc
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32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	32.7	33.0	33.0
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H	4	4	4	N	4	4	4	4	4	4	4	4	4	4	4	4	4
US-08-493-092-3	US-08-952-014C-1	US-08-952-127-1	US-08-642-274D-1	US-08-629-001A-1	US-09-528-784A-45	US-09-528-784A-7	US-08-723-142A-45	US-08-723-142A-7	US-08-990-571-45	US-08-990-571-7	US-08-845-258-45	US-08-845-258-7	US-09-134-001C-567	US-09-134-001C-2388	US-09-134-001C-458	US-09-470-512A-3	US-09-350-600-36
Sequence 3, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 1, Appli	Sequence 45, Appl	Sequence 7, Appli	Sequence 45, Appl	Sequence 7, Appli	Sequence 45, Appl		Sequence 45, Appl	Sequence 7, Appli	Sequence 567, A	Sequence 2388,	Sequence 458, A	Sequence 3, Appl	Sequence 36, App

ALIGNMENTS

APPLICANT: Chen, Bin TITLE OF INVENTION: 5(Upstream Region Seque TITLE OF INVENTION: and Uses Thereof FILE REFERENCE: D6015 CURRENT APPLICATION NUMBER: US/09/193,792B CURRENT FILLING DATE: 1998-11-17 PRIOR APPLICATION NUMBER: US 60/065,113 PRIOR FILLING DATE: 1997-11-18 NUMBER OF SEQ ID NOS: 20 SEQ ID NO 20 ; OTHER INFORMATION: PstI-EheI fragment (PE1.3) inserted in the multiple; OTHER INFORMATION: cloning region of a pGLe enhancer vector to construct; OTHER INFORMATION: the transfection plasmid US-09-193-792-20 Sequence 20, Application US/09193792B Patent No. 6180344 GENERAL INFORMATION: 5(Upstream Region Sequences and Uses Thereof 35.2%; 64.7%; Score 22.2; D: Pred. No. 9.4; 0; Mismatches 0; BB 18; 4. of the MYOD1 Gene Length 1312; Indels 0 Gaps 0;

HIGH AFFINITY RECEPTOR-LIKE

PROTEIN IMMUNOGLOBULIN

COMPUTER READABLE FORM

MEDIUM TYPE: Dis COMPUTER: IBM CO OPERATING SYSTEM:

IBM Compatible

Diskette

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US-09-213-389-2
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NAME: Billings, Lucy J.

REGISTRATION NUMBER: 36,749
REFERENCE/DOCKET NUMBER: PF-0
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO:
                              CLASSIFICATION:
PRIOR APPLICATION NUMBER:
                                                                                                          SOFTWARE: FastSEQ for CURRENT APPLICATION DATA:
                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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LIBRARY: BRAIN
CLONE: 927955
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: FastSEQ for Windows Version CURRENT APPLICATION DATA:
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FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
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                                                                                                                                       COMPUTER: IBM COMPONENTING SYSTEM:
                                                                                                                                                                                                                    COUNTRY:
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                                 08/916,902
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Pred. No. 10;
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; OTHER INFORMATION: PstI fragment (PP2.0) used as a hybridization probe
US-09-193-792-2
                                                                                                                                                                                                                                                    us-09-193-792-1/c
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                                                                                                                                                                                             Sequence 1, Application US/09193792B Patent No. 6180344 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 33; Conserv
NUMBER OF SEQ ID NOS:
SEQ ID NO 1
LENGTH: 2284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQ ID NO 2
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                                                   CURRENT APPLICATION NUMBER: US/09/193,792B
CURRENT FILING DATE: 1998-11-17
PRIOR APPLICATION NUMBER: US 60/065,113
PRIOR FILING DATE: 1997-11-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/193,792B CURRENT FILING DATE: 1998-11-17 PRIOR APPLICATION NUMBER: US 60/065,113 PRIOR FILING DATE: 1997-11-18 NUMBER OF SEQ ID NOS: 20
                                                                                                                            APPLICANT: Chen, Bin
TITLE OF INVENTION: 5( Upstream Region Sequences of the MYOD1 Gene
TITLE OF INVENTION: and Uses Thereof
FILE REFERENCE: D6015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: 5( Upstream Region Sequences TITLE OF INVENTION: and Uses Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                               1091 AGCCCTAGGCGGCTGACACTTGGCTCTCCGGCACGCCCTTTCCAAACCTCT 1041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IMMEDIATE SOURCE:
LIBRARY: BRAIN
CLONE: 927955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS LENGTH: 1669 base pair
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: TELEPHONE: 415-855-0555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 435 GGAAGTAACCCTATTTCCGTGTATATCGGGTACACAATTTGGGGGGTCAGTAATGTTTAT 493
                                                                                                                                                                                                                                                                                                                                                              11 AACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCCAAGCTTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 GAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTTTAT 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2021
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1669 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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64.7%;
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Pred. No. 1
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Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 2021;
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; FEATURE:
; NAME/KEY: promoter
; LOCATION: -1537..747
; OTHER INFORMATION: 5( upstream promoter region of the human MYOD1 gene
US-09-193-792-1
US-09-605-785-535/c
Sequence 535, Application US/09605785; Patent NO. 6321716; GENERAL INFORMATION:
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US-08-961-527-158/c
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                                                                RESULT 7
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Best Local Similarity
Matches 33; Conserv
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                 TELEFAX: (301) 309-85
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Diskette,
                                                                                                              1897 CGAATCATAGAATTATTCGGTTGAAACTTGGTGACGAAGTATGGGGATTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 424
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: MSDO
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1210 AGCCCTAGGCGGCTGACACTTGGCTCTCCGGCACGCCCTTTCCAAACCTCT 1160
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                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: PE
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Brookes, A. Anders
REGISTRATION NUMBER: 36,373
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 9410 Ke
CITY: Rockville
                                                                                                                                                                                                                                                         TOPOLOGY:
                                                                                                                                                                                                                                                                                                      LENGTH:
                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
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                                                                                                                                         3 CGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCC 53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  158, Application US/08961527
5. 6420135
                                                                                                                                                                                                                                                                                 i: 3140 base pairs nucleic acid
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9410 Key West Avenue
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                                                                                                                                                                            Conservative
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64.7%;
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                                                                                                                                                                        Score 22.2; D
Pred. No. 12;
0; Mismatches
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Pred. No. 11;
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; SOFTWARE: FastSEQ for Wir
; SEQ ID NO 535
; LENGTH: 6082
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-605-785-535
                                                         ; LENGTH: 6082
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-439-313-535
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                                                                                                                 SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 535
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Best Local Similarity
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Best Local Similarity
                                                                                                                                            CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
NUMBER OF SEQ ID NOS: 575
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                                                                                                                                                                                       APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER FILE REFERENCE: 210121.427C16
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                                                                                                                                                                                                                                                                                                                                                                                                              tent No.
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                                                                                                                                                                                                                                                                                                                                                                                                              535, Application US/09439313
5. 6329505
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Dillon, Davin C
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                                                                                                                                                                                                                                                                                                        Jiang Yugui
Reed, Steven G.
                                                                                                                                                                                                                                                            Fanger, Gary
Retter, Mark
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                                                                                                                                                                                                                                                                                            Kalos, Michael
                                                                                                                                                                                                                                                                                                                                     Harlocker,
                                                                                                                                                                                                                                                                                                                                                      Mitcham, Jennifer L.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Carter, Darrick
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kalos, Michael
 Conservative
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ick, Thomas S.
                                                                                                                                                                                                                                                   John
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2000-06-27
               34.6%;
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Score 21.8; Di
Pred. No. 19;
0; Mismatches
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Pred. No. 19;
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   22;
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AAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTTTA 62

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2431 ACAGTGACATTTAGCATACTTTGTTTGCCCCAGTATGAAAGCCACCAATCTTGA 2375

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; LOCATION: (4535)
; OTHER INFORMATION: n-A,T,C
US-09-605-785-536
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                                                                                                                                                                                                                                                                                              US-09-439-313-536/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 536
LENGTH: 6140
                                                                                                                                                                                                                                                                Sequence 536, Application US/09439313. Patent No. 6329505
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APPLICANT: Day, Craig
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C9
CURRENT APPLICATION NUMBER: US/09/439,313
CURRENT FILING DATE: 1999-11-12
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APPLICANT: Hepler, William
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF PROSTATE CANCER
FILE REFERENCE: 210121.427C16
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ORGANISM: Homo sapiens
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                                                                                                                                                     Jiang Yuqui
Reed, Steven G.
                                                                                                                                                                                              Dillon, Davin C.
Mitcham, Jennifer L.
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Kalos, Michael D.
                                                                                                          Fanger, Gary
Retter, Mark
                                                                                                                                        Kalos, Michael
                                                                                        Solk, John
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Pred. No. 20;
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US-08-727-118-1
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; LLCATION: (4535)
; OTHER INFORMATION: n-A,T,C or
US-09-439-313-536
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SOFTWARE: FASTS
SEQ ID NO 536
FENGTH: 6140
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Best Local Similarity
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Best Local Similarity
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SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                            TELEFAX: (617) 248-710
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: VITO PHD, CHRISTINE C.
REGISTRATION NUMBER: 39,061
REFERENCE/DOCKET NUMBER: CRP
                                                                                                                                                         FEATURE:
                                                                                                                                                                      MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: MIYAZONO, KOHEI APPLICANT: SAMPATH, KUBER T. APPLICANT: COHEN, CHARLES M.
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                                                                                                                                                                                      TOPOLOGY:
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                                                                                                                                                                                                 STRANDEDNESS: single
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2 GCGGAAAGTAACCTTATGCATTTGTAACCTTGCTTACCCGGTATGCCGGTTCCCAAGCT 59
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3ER: CRP-121 [2054/91]
                                             Score 21.2; D
Pred. No. 30;
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                                                                                          APPLICANT: Reynolds
                                                                                                                           Sequence 1, Application US/08680506C Patent No. 6008013
                                                                                                                                                                                                                                                                                                                  Query Match
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Patent No. 600801
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Best Local Similarity 70.0%;
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CURRENT APPLICATION NUMBER: US/08/680,506C
CURRENT FILING DATE: 1996-07-08
EARLIER APPLICATION NUMBER: 60/021,672
EARLIER FILING DATE: 1996-07-05
                                                                           APPLICANT: Reynolds, Paul R.
TITLE OF INVENTION: CHONDROCYTE PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/08/680,506C CURRENT FILING DATE: 1996-07-08
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TITLE OF INVENTION: CHONDROCYTE PROTEINS
FILE REFERENCE: 176/60091
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NUMBER OF SEQ ID NOS: 18
                                                            FILE REFERENCE: 176/60091
                                                                                                                                                                                                                                                                                                                                                             LENGTH: 5027
TYPE: DNA
ORGANISM: Gallus gallus
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TYPE: DNA
ORGANISM: Gallus gallus
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TITLE OF INVENTION: CHONDROCYTE PROTEINS
FILE REFERENCE: 176/60091
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Pred. No. 46
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Pred. No. 3
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US-08-181-271A-36/c
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Best Local :
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                                                                                                                                                                                                                                                                                OPERATING SYSTEM: PC-D
SOFTWARE: PatentIn Rel
CURRENT APPLICATION DATA:
                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 1-APR-1991
                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                           FILING DATE: 6-FEB-1989 PRIOR APPLICATION DATA:
                                                                      PRIOR APPLICATION DATA:
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                                                                                                                                        APPLICATION NUMBER: US 07/937,197 FILING DATE: 6-NOV-1992
                                                                                                                                                                                      APPLICATION NUMBER: US 0 FILING DATE: 16-JUL-1993
                                                                                                                                                                                                                                                  APPLICATION NUMBER: US FILING DATE: 13-JAN-94
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ZIP: 10532
                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                       CLASSIFICATION:
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sperison, Jeffrey)
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SYSTEM: PC-DOS/MS-DOS
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              OS 07/165,667
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Pred. No. 5;
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Search completed: May 24, 2003, 15:37:41 Job time : 32.0806 secs
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ORIGINAL SOURCE:

ORGANISM: Cucumis sativus

INDIVIDUAL ISOLATE: Cucumber Chitinase Genomic DNA

IMMEDIATE SOURCE:

CLOME: pBSCucchrcht5

US-08-181-271A-36
                                                                                                                                                               Query Match 33.0%;
Best Local Similarity 70.0%;
Matches 28; Conservative
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APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/768,122
APPLICATION NUMBER: US 07/768,122
APPLICATION DATA:
APPLICATION DATA:
APPLICATION UNBER: US 07/580,431
APPLICATION UNBER: US 07/368,672
APPLICATION UNBER: US 07/368,672
APPLICATION UNBER: US 07/368,672
APPLICATION UNBER: US 07/329,018
APPLICATION UNBER: US 07/329,018
APPLICATION UNBER: US 07/329,018
APPLICATION UNBER: US 08/045,957
ATTORNEY/AGENT INFORMATION:
NAME: Elmer, James Scott
REFERENCE/DOCKET NUMBER: S-19825/P1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919)541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (919)541-8614
TELEPAX: (919)541-8689
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 13124 base pairs
                                                                              8248 TTGGAACTTGGTGGACGGGGCTTCCGGTTGAGAAGCTGTA 8209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US 07/632,441
FILING DATE: 21-DEC-1990
PRIOR APPLICATION UNMBER: US 07/425,504
FILING DATE: 20-OCT 1989
PRIOR APPLICATION UNMBER: US 07/425,504
PRIOR APPLICATION DATA:
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APPLICATION NUMBER: US (
FILING DATE: 6-APR-1993
                                                                                                                   23 TTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTTTA 62
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Pred. No. 57;
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Maximum Match 100%
Listing first 45 summaries
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Sequence 11, Appl sequence 5, Appl Sequence 7, Appl Sequence 7, Appl Sequence 6, Appl Sequence 11, Appl Sequence 21, Appl Sequence 23, Appl Sequence 25, Appl Sequence 29, Appl Sequence 20, Appl Sequence 20, Appl Sequence 21, Appl Sequence 21, Appl Sequence 22, Appl Sequence 21, Appl Sequence 2148, App Sequence 148, A
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octuence not who	2	Sequence 536, App	Sequence 536, App	Sequence 536, App	Sequence 535, App	Seguence 535, App	e 535,	535,	Sequence 535, App	Sequence 535, App	Sequence 535, App	Sequence 1007, Ap	Sequence 1006, Ap	Sequence 613, App	3503	Sequence 3832, Ap	Sequence 293, App	Sequence 1, Appli	Sequence 227, App	Sequence 21, Appl	Sequence 2855, Ap	Sequence 3, Appli	Sequence 3369, Ap	Sequence 5603, Ap	Sequence 762, App	Sequence 646, App

THENSENTS

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RESULT 1
US-09-963-803-11
; Sequence 11, Application US/09963803
; Publication No. US20030028922A1
; GENERAL INFORMATION:
RESULT 2
US-09-963-803-3
; Sequence 3, Application US/09963803
                                                                                                                                                                                                                                                                                                                                                                      US-09-963-803-11
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Best Local S
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ. ID NOS: 39
NUMBER OF SEQ. ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence FEATURE: OTHER INFORMATION: Directional OTHER INFORMATION: S4
                                                                                                             61
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                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                  1 GGCGGAAAGTAACCTTATGCATTTTGTAACCTTGGTTACCCGGTATGCCCGGTTCCCCAAGCTT 60
                                                                                                           TAT
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CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTMARE: PRIENTIN VERSION 3.1
SEQ ID NO 3
LENGTH: 317
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; LOCATION: (1)..(317)
; OTHER INFORMATION:
US-09-963-803-3
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LENGTH: 371
                                       Matches
                                                                        Query Match
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                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
NUMBER OF SEQ ID NOS: 39
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava velo mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
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                                                                                                                          NAME/KEY: promoter LOCATION: (1)..(371) OTHER INFORMATION:
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OTHER INFORMATION: Promoter MPr1116
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                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                           TYPE: DNA
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                                                       Local Similarity
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100.0%; Pred. No. 1.9e-1;
tive 0; Mismatches (
                                     0
                                                    Score 63;
Pred. No.
                                   Mismatches
                           1.9e-15;
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US-09-963-803-4
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SEQ ID NO 7
LENGTH: 301.
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                                                                                  SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 348
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Best Local :
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APPLICANT: Meristem Therapuetics
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PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                                                                                    APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
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CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05
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                                                                     TYPE: DNA
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OTHER INFORMATION: promoter MPr1117
                      FEATURE:
                                          ORGANISM: Artificial Sequence
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nes 62; Conserv
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Pred. No. 8.1e-15;
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RESULT 7
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; LOCATION: (1)..(398)
; OTHER INFORMATION:
US-09-963-803-6
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GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
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Best Local :
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Best Local Similarity
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
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NAME/KEY: promoter
LOCATION: (1)..(348)
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION NUMBER: PCT 1B00/00370 PRIOR FILING DATE: 2000-10-05 NUMBER OF SEQ ID NOS: 39 SOFTWARE: Patentin version 3.1
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
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OTHER INFORMATION:
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TYPE: DNA
ORGANISM: Artificial Sequence
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Pred. No. 8.9e-15;
0; Mismatches 1;
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Pred. No. 8.5e-15;
0; Mismatches 1.
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PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 243
TYPE: DNA
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Best Local (
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LENGTH: 392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                  Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello TITLE OF INVENTION: Virus and cassava vein mosaic virus FILE REFERENCE: 18433042 CURRENT APPLICATION NUMBER: US/09/963,803 CURRENT FILING DATE: 2001-09-26 CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Meristem Therapuetics
                                                                                                                                                                                                                                       NAME/KEY: promoter LOCATION: (1)...(392) OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: promoter LOCATION: (1)..(243) OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: promoter MPrl164
                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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257 CA 258
                                                                          197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        153 TTTGTAACTTGGTTACCCGGTATGCCGGTTCCCCAAGCTTTAT 194
                                     61 TA 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22 TTTGTAACTTGGTTACCCGGTATGCCGGTTCCCAAGCTTTAT 63
                                                                                                                                                                   Local Similarity
                                                                                         1 GGCGGAAAGTAACCTTATGCATTTTGTAACCTTGGTTACCCGGTATGCCCGGTTCCCAAGCTT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                        GGCGGAAAGTAACCTTATGCATTTGTAATTTGGTTACGTGGCATGCGTCATCCCTTACGT 256
                                                                                                                                                49;
                                                                                                                                                  Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    243 bp Fragment from the intergenic region of commelina yellow ttle virus % \left( 1\right) =\left\{ 1\right\} 
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                                                                                                                                                               65.48;
79.08;
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                                                                                                                                                Score 41.2; DB
Pred. No. 1e-06;
0; Mismatches
                                                                                                                                                                                 DB 9;
                                                                                                                                                  13;
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US-09-963-803-23 ; Sequence 23, Application US/09963803 ; Publication No. US20030028922A1

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                                                                                                                                                  ; NAME/KEY: promoter
; LOCATION: (1)..(472)
; OTHER INFORMATION:
US-09-963-803-25
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                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 25
LENGTH: 472
                                                                          Matches
                                                                                                               Query Match
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SEQ ID NO 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow TITLE OF INVENTION: virus and cassava vein mosaic virus FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR PPLICATION NUMBER: PCT 1B00/00370
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 3000-10-05
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                                                                                                                                                                                                                               FEATURE:
                                                                                                                                                                                                                                              OTHER INFORMATION: promoter MPr1169
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                                                                                                                                                                                                                                                                                                       TYPE: DNA
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OTHER INFORMATION: promoter MPr1167
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 278
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                 1 GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC 37
                                                                                              Local
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GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC 314
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                                                                                              Similarity
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                                                                          Conservative
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                                                                                          58.7%;
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95.2%;
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                                                                                            Score 37;
Pred. No.
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                                                                        Mismatches
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                                                                                                             DB 9; Length 472;
                                                                                            5e-05;
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; LOCATION: (1)..(54:
; OTHER INFORMATION:
US-09-963-803-24
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                                                                                                                           US-09-963-803-19
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                                                                                                                                                                                                                                                                                           SEQ ID NO 19
LENGTH: 39:
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                                                      Matches
                                                                                    Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILLING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS:
SOFTWARE: PatentIn ver
                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PRIOR FILING DATE: 2000-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: virus and cassava vein mosaic virus
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SOFTWARE: PatentIn ver
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE:
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                                                                                                                                         NAME/KEY: promoter LOCATION: (1)..(39) OTHER INFORMATION:
                                                                                                                                                                                                    OTHER INFORMATION: promoter MPrl162 FEATURE:
                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                     Local Similarity
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1 GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTAC 37
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37; Conserv
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                                                  Conservative
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                                                                                                                                                                (393)
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                                                                                                                                                                                                                                                                                                                                                                     2000-10-05
                                                                   56.2%;
97.3%;
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                                                  0,
                                                Score 35.4; DB 9;
Pred. No. 0.0002;
0; Mismatches 1;
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Pred. No. 5.2e-05;
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FEATURE:
NAME/KBY promoter
LOCATION: (1)..(600)
OTHER INFORMATION:
US-09-963-803-22
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; LOCATION: (1)..(462)
; OTHER INFORMATION:
US-09-963-803-20
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Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
Query Match
Best Local Similarity 97.3
Matches 36; Conservative
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Best Local Similarity
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR PPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 300
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn version 3.1
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PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                              LENGTH: 600
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 462
TYPE: DNA
ORGANISM: Artificial Sequence
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OTHER INFORMATION: promoter MPr1163
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97.3%;
                      56.2%;
97.3%;
Score 35.4; DB 9;
Pred. No. 0.00024;
0; Mismatches 1
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Pred. No. 0.00022;
0; Mismatches 1
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                                      9;
                                      Length 600;
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  Indels
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  0;
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CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 2913
LENGTH: 434
TYPE: DNA
ORGANISM: Homo sapien
US-09-867-701-2913
Search completed: May 24, 2003, 15:36:01 Job time : 51.1129 secs
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APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
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ALIGNMENTS

VERSION
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ORGANISM RESULT 1 C88100 LOCUS JOURNAL COMMENT FEATURES REFERENCE ACCESSION DEFINITION TITLE AUTHORS source Mus musculus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Sciurognathi; Muridae; Murinae; Mus 1 (bases 1 to 501) 1 C88100 Mouse fertilized clone J0256H01 3', mRNA C88100 Unpublished (1998)
Contact: Hirofumi Doi
Doi Bloasymmetry Project, ERATO
Doi Bloasymmetry Project, ERATO
Japan Science and Technology Corporation (JST)
WBG Marive East 12F, 2-6 Nakase, Mihama-ku, Chiba 261-71, Japan
Email: hd@bioa.jst.go.jp.
Location/Qualifiers Systematic analyses of genes expressed in fertilized (The ERATO/Doi Project at Wayne State University) C88100.1 GI:2920057 Doi, H. house mouse /organism="Mus musculus" /strain="C57BL/6J" /db_xref="taxon:10090" 501 bp mRNA linear EST 11-MAR one-cell-embryo cDNA Mus musculus cDNA sequence. Paonessa, P.D. and EST 11-MAR-1998 mouse eggs

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1 (bases 1 to 501)

Mahairas,G.G., Wallace,J.C., Smith,K., Swartzell,S., Holzman,T., Keller,A., Shaker,R., Furlong,J., Young,J., Zhao,S., Adams,M.D. a.
                                                                                                                                                                                                                                                                                          Email: jwallace@u.washington.edu
Sequence Tagged Connector
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h Throughput Sequencing Center
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/note="Organ: mammary gland; Vector: pT7T3D-Pac (Pharmacia ) with a modified polylinker; Site_1: Not I; Site_2: Eco RI; 1st strand cDNA was primed with a Not I - oligo(dT) primer [5'
                                                                                                                            /note="Organ: sperm; Vector: pBeloBAC11; BAC
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Whitton,C., Daub,J., Ware,J., Quail,M., Hall,N., Barrell,B., Foster,J., Guiliano,D., Slatko,B. and Blaxter,M.
Genome survey sequences from the human parasitic nematode Brugia
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Fax: +44 131 670 5450
Email: mark.blaxter@ed.ac.uk
Sequenced from the Brugia malayi BAC library constructed by Claire
Whitton and Dr Mike Quail. The sequence was generated by The
Pathogen Sequencing Unit, The Sanger Institute, Cambridge, UK in
collaboration with Mark Blaxter, ICAPB, University of Edinburgh,
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Institute of Cell, Animal and Population Biology
University of Edinburgh
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/sex="Mixed (male and female)"
/tissue_type="whole parasite"
/dev_stage="microfilaria (LL)"
/note="Vector: pBACe3 6; Site_1: BamH I; Brugia malayi
/fote="Vector: pBACe3 6; Site_1: Ba
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/strain="TRS"
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Eukaryota; Metazoa; Platyhelminthes; Trematoda; Digenea;
Strigeidida; Schistosomatoidea; Schistosomatidae; Schistosoma.

1 (bases 1 to 835)
1 (Passier, M.C., Pierce, R.J., Merlin, F., Hirai, H., Wu, W.,
Williams, D.L., Johnston, D., Loverde, P.T. and Le Paslier, D.
Construction and characterization of a Schistosoma mansoni
Dacterial artificial chromosome library
                                                                                                                                                                                                                                                   CNS070W0 835 bp DNA linear GSS 10-OCT-2001 T7 end of clone 024AH05 of library SmBAC1 from strain Puerto-Rican of Schistosoma manson1, genomic survey sequence.
                                                                                                                                                                                                   GSS.
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Genomics 65 (2), 87-94 (2000)
20247247
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Seq primer: M13 -21 primer
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CNS sequencing ID=DGOAA024AH05CP1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Institut Pasteur de Lille
1 rue du Professeur A. Calmette, 59019-Lille,
Tel: (33) (0)3 20877783
Fax: (33) (0)3 20877888
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Contact: Pierce RJ
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/lab_host="Blomphalaria glabrata"
/lab_host="Blomphalaria glabrata"
/note="vector: pBeloBAC 11; Site_1: Hind III; Partially
/note="vector: pBeloBAC 11; Site_1: Hind III digested and size-selected S. mansoni cercarial
DNA was ligated into Hind III digested pBeloBAC 11 vector
and used to transform E. coil pBlOB. The complete library
contains 23808 clones from 4 independent
sizing-ligation-transformations. Average insert size
ranges from 70-127 kb and genome coverage is 7.9-fold."
a 141 g 261 t 13 others
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/strain="Puerto-Rican"
/db_xref="taxon:6183"
/clone="024009"
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Partially Hind III digested and size-selected S. mansoni cercarial DNA was ligated into Hind III digested pBeloBAC 11 vector and used to transform E. coli DH10B. The complete library contains 23008 clones from 4 independent sizing-ligation-transformations. Average insert size ranges from 70-127 kb and genome coverage is 7.9-fold.
                                                                                                                                                                                                                                        Email: est@watson.wustl.edu
Insert Size: 721
High quality sequence stops: 315 Source: IMAGE Consortium, LLNL
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Insert Length: 721 Std Error: 0.00
Seg primer: M13RP1
Seg primer: M13RP1
                                                                                                                                                                                                                                                                                                                                                                                                                        Washington University Schwa4444 Forest Park Parkway, Tel: 314 286 1800 Fax: 314 286 1810
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Hilller,L., Clark,N., Dubuque,T., Elliston,R., Hawkins,M., Holman,M., Kucaba,T., Le,M., Lennon,G., Marra,M., Parsons,J.,Rifkin,L., Rohlfing,T., Soares,M., Tan,F., Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P. and Wilson,R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    426 bp mRNA lir
Yj74d07.rl Soares breast 2NbHBst Homo sapiens
IMAGE:154477 5', mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1995)
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/strain="puerto-Rican"
/db_xref="taxon:6183"
/clone="024AH05"
/clone_lib="SmBAC1"
/note="end : T7"
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                                                   /clone_lib="Soares breast 2NbHBst"
                                                                        /db_xref="taxon:9606"
/clone="IMAGE:154477"
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'dev_stage="adult"
                            'sex="Female"
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                                                                                                                                                                                                                                                                   Email: est@watson.wustl.edu
This clone is available royalty-free through LLNL; contact the
IMAGE Consortium (info@image.llnl.gov) for further information.
Seq primer: -28ml3 rev2 ET from Amersham
High quality sequence stop: 338.
                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 434)
Hiller,L., Allen,M., Bowles,L., Dubuque,T., Geisel,G., Jost,S.,
Kucaba,T., Lacy,W., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B.,
Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,
,T., Waterston,R. and Wilson,R.
Washu-Merck EST project 1997
Unpublished (1997)
                                                                                                                                                                                                                                                                                                                                                                Washington University School of Medicine
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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/db_xref="taxon:9606"
/clone="IMAGE:724752"
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BG150109 519 bp mRNA linear EST 05-FEB-2001 nad51g04.x1 NCI_CGAP_Lu24 Homo sapiens cDNA clone IMAGE:3407047 3', mRNA sequence.
BG150109
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1 (bases 1 to 493)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anaty
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zs44d05.rl NCI_CGAP_GCB1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert Strausberg, I
Email: cgapbs-r@mail.nih.gov
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Unpublished (1997)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Contact: KOUDEL CONTACT IN THE STATE OF THE 
                                                                                                                               Tumor Gene Index
Unpublished (1997)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                       NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 554)
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similar to contains Alu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AI970488
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High quality sequence stop: 483.
Location/Qualifiers
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Tissue Procurement: Christopher Moskaluk, M.D., Ph.D.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Unpublished (1997)
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1 (bases 1 to 519)
NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Emmert-Buck, M.D., Ph.D.
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CDNA Library Arrayed by: Greg Lennon, Ph.D.
DNA Sequencing by: Washington University Genome Sequencing Center
Clone distribution: NCI-CGAP clone distribution information can be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Soares and M. Fatima Bonaldo.
106 c 81 g 152 t
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/db_xref="taxon:9606"
/clone="IMAGE:3407047"
/clone_lib="NCI_CGAP_Lu24"
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Pred. No. 41;
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                                                   Moskaluk, M.D.,
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RESULT 11
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Best Local
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                                                                                                                                                                                                                                               Unpublished (2007),
Contact: John Quackenbush
The Institute for Genomic Research
The Institute for Genomic Research
                                                                                                                                                                        Plate: 211
                                                                                                                                                                                                                  9712 Medical Center Dr.,
Tel: 301 838 3528
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                  Assessment of gene expression patterns in a model of colon tumor metastasis using a 19,200 element cDNA microarray Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                  Hegde, P., Qi, R., Abernathy, K., Dharap, S., Gaspard, R., Gay, C., Holt, I.E., Saeed, A.I., Sharov, V., Lee, N.H., Yeatman, T.J. and
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Seg primer: -40UP from Gibco
High quality sequence stop: 457.
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                                                                                                                                                                                                                                                                                                                                                                                   Quackenbush,
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Mammalia; Eutheria; Primates;
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Similarity 63.3%;
38; Conservative
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                  255
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          /organism="Homo sapiens"
/ob_xref="taxon:9606"
/db_xref="taxon:9606"
/clone_lib="MAGE resequences, Note="Vector: pBluescriptSKm"
/note="Vector: pBluescriptSKm"
135 c 112 g 228 t
                                                                                                                   Location/Qualifiers
1. .730
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polylinker; Site_1: Not I; Site_2: Eco RI; Plasmid DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /tissue_type="pooled germ cell tumors"
/lab_host="DH10B"
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/clone="IMAGE:2479616"
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Pred. No. 4
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Mismatches
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ens cDNA,
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                                                                                                                                          GGTAAGCAACATTGTTCATTTGTTACTGGGTGTTCTAGATCAAACCTTCACAAGCTATAT 335
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                               BF970066 918
602272333F1 NIH_MGC_84 Homo :
mRNA sequence.
BF970066
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               http://image.llnl.gov
Plate: LLCM866 row: e column: 19
High quality sequence stop: 540.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CDNA Library Preparation: CLONTECH Laboratories, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONTECH Laboratories, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 (bases 1 to 811)
NIH-MGC http://mgc.ncl.nih.gov/.
National Institutes of Health, Mammalian
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BF107114.1 GI:10889555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 mRNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BF107114
                                                                                                                                                                                                                                     Similarity
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                                                                                                                                                                                                                                                                                                            240
                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                             /clone_lib="NIH_MCC_79"

/clone_lib="NIH_MCC_79"

/lab_host="DH10B (T] phage-resistant)"

/note="Organ: placents; Vector: pDNR-LIB (Clontech);

// is and 3 daptor sequence: 5'-ACGGCCATTATGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGACATGTGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3'

/where B = A, C, or G and N = A, C, G, or T). Average insert size 1.3 kb (range 0.5-4.0 kb). 15/15 colonies contained inserts by FCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo Alto, CA). Note: this is a NIH_MGC

// ith__..."
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                                                                                                                                                                                                                                                                                                      Library."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                               39.48;
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Pred. No. 47;
0; Mismatches
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Pred. No. 46;
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                               mRNA linear EST 22-Ja
s cDNA clone IMAGE:4360233
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AZ315498/c
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KEYWORDS
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TITLE
JOURNAL
                                                                                                                       JOURNAL
                                                                                                                                                          TITLE
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                                                                                                                                                                                                                                 AUTHORS
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VERSION
KEYWORDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 GGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCCAAGCTTTAT 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                   A2315498 487 bp D 1M0032D15R Mouse 10kb plasmid UUGC1M Clone UUGC1M0032D15 R, DNA sequence. A2315498 GI:10362388
                                                                                                                                                                                                                                                                                                                                                                                                                                   GSS.
                                                                                      University of Utah
University of Utah
                                                                                                                                     Contact: Robert B.
                                                                                                                                                          Unpublished (2000)
                                                                                                                                                                                 plasmid inserts
                                                                                                                                                                                                       Mouse whole genome
                                                                                                                                                                                                                              and Wright, D., Weiss, R.
                                                                                                                                                                                                                                            Dunn,D., Aoyagi,A., Barber,M., Beacorn,T., Duval,B., Hamil,C., Islam,H., Longacre,S., Mahmoud,M., Meenen,E., Pedersen,T., Reilly, Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
1 (bases 1 to 487)
                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Robert Strausberg, Ph.D. Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                              house mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  http://image.llnl.gov
Plate: LLAM10001 row: b column:
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National Institutes of Health, Mammalian
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1 (bases 1 to 918)
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EST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CDNA Library Preparation: Life Technologies, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
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801 585 5606
801 585 7177
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                                                                   Biomedical
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/lab_host="DH10B (phage-resistant)"
/note="Organ: adrenal gland; Vector: pCMV-SPORT6; Site_1:
NotI; Site_2: SalI; Cloned unidirectionally; oligo-dT
primed. Average insert size 1.229 kb. Library enriched for
full-length clones and constructed by Life Technologies.
Note: this is a NIH_MGC Library."
171 c 173 g 305 t
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/clone_lib="NIH_MGC_84"
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/db_xref-"taxon:9606"
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                                                                                                              Genome Center
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                                                                 Polymers Research Bldg.,
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Pred. No. 49;
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smid UUGClM library Mus musculus
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                                                                 2030 E.,
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ORGANISM
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BH116178/c
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ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DEFINITION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Focus
                                                                                                                               TITLE
JOURNAL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           190 GGCTGAGATTATTTTCCTGCTGTTCAAACCCTGTTACCCAGTATGGTGGCTACAAAGAGT 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 130 TCT 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 TAT 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GGCGGAAAGTAACCTTATGCATTTGTAACTTGGTTACCCGGTATGCCCGGTTCCCAAGCTT 60
Zhao,S., Nierman,W., Malek,J., Shatsman,S., Akin Tsegaye,G., Geer,K., Krol,M., Shvartsbeyn,A., Ge Russell,D., de Jong,P. and Fraser,C.M.
Mouse BAC End Sequences from Library RPCI-24
Unpublished (1999)
Other.GSS: RPCI-24-223H11.TJ
Contact: Shaying Zhao
Department of Eukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD 20850, US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            39;
                                                                                                                                                                                                                                                                                                                                                                                                                     , DNA sequence.
BH116178
                                                                                                                                                                                                                                                                                                                                                                                                                                                      BH116178 508 bp DNA LINE RPCI-24-223H11.TV RPCI-24 Mus musculus genomic
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Location/Qualifiers
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                                                                                                                                                                                                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                      Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                              BH116178.1 GI:14956155
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                                                                                                                                                                                                                                                Mammalia; Eutheria;
1 (bases 1 to 508)
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                                                                                                                                                                                                                                                                                                                                                 nouse mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Insert Length: 10000 Std Erro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                129
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pWD42 (gil4732114|gb]AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /lab_host="E. Coli strain XL10-Gold, T1-resistant, F-"/note="Vector: PMD42nv; Purified genomic DNA from M. musculus C57BL/60 (male) was obtained from the Jackson Laboratory Mouse DNA Resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /db_xref="taxon:10090"
/clone="UUGC1M0032D15"
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/strain="C57BL/6J"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /sex="Male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Mouse 10kb plasmid UUGC1M library"
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61.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   49;
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199 TTATTCATTTCTAATTTTGTTAACTTGTATGCTGTCTCAGAGCCTTT 15:
                     15 TTATGCATTTGTAACTTGGTTACCCGGTATGCCGGTTCCCCAAGCTTT 61
                                                                                  33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Class: BAC ends.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clones are derived from the mouse BAC library RPCI-24. For BAC library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC Resources (http://www.chori.org/bacpac/orderingframe.htm). BAC end page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html Plate: 223 row: H column: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Email: szhao@tigr.org
                                                                                                      Similarity
                                                                                                                                                                                         220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              primer: T7
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301 838 0208
                                                                                  Conservative
                                                                                                                                                                                                                         /cell_type="Spleen/Brain"
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
RPCI-24 Mouse BAC Library produced by Pieter de Jong. The
library was cloned in the pTARBAC1 cloning vector at the
BamH1 sites using MboI partially digested male C57BL/6J
                                                                                                                                                                                                            DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                       /organism="Mus musculus"
/strain="C57BL/6J"
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/clone="RPCI-24-223H11"
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70.2%;
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Pred. No. 49;
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Search completed: May 24, 2003, 15:33:24 Job time: 831.806 secs

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GenCore version 5.1.4_p5_45

Database : Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries Minimum DB seq length: 0
Maximum DB seq length: 2000000000 Total number of hits satisfying chosen parameters: Searched: Title: Perfect score: OM nucleic - nucleic search, using sw model Scoring table: Sequence: Run on: US-09-963-803-13 2054640 seqs, 14551402878 residues IDENTITY_NUC Gapop 10.0 , Gapext 1.0 May 24, 2003, 10:21:51; Search time 350.903 Seconds (without alignments) 5225.021 Million cell updates/sec 1 catgctgcagactagtggat.....aagggatgacgcatgccact 63 GenEmbl:* GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd. 9b_pat: 9b_ph: 9b_pr: 9b_pr: 9b_ro: 9b_sts: 9b_sts: 9b_un: 9b_un: em_mu:*
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gb_in:* em_htgo_hum:*
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em_sy:* em_hum:*
em_in:* em_ba:* em_fun:* gb_om: * 4109280

Pred. No. is the number of results predicted by chance to have a

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

RESULT 1 AX036610 LOCUS DEFINITION ACCESSION VERSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS JOURNAL FEATURES	44.55	44 43 43	4 3 9 0 1 0 9 0	37 38	3 33 33 75 A	c 32	30 31	28 29		, C 24	c 22	20 21	18	16 17	c 14 15	13	11	1 9	8 7	100	Α . π	ω κ) L	Result
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om Patent FR2791358. :11226205 truct truct truct uences. 63) 63) 820, M. and Gruber, V. 1358-A 16 29-SEP-2000; PEUTICS (FR) ion/Qualifiers	AR152425 ALIGNMENTS	AR110594 AR150993 AR157416	AX026717 SYNCAMVCM5	7 N #	79	ARCAMVPR SCO308514	E01311 I04847	AX405116 BD001990	AX033493 AX044092	AF434748 AF434750	AF434747 AF434746	AF434749 AX164073	AX093047 AX093052	AX036754 AX036756	AX036755 AX036753	AX036603 AX036739	AX036616	AX036611	AX036758 AX036757	AX036759	AX036741	AX036601	3661	ID
linear PAT 16-NOV-2000	0, ,	AR403123 sequence AR110594 sequence AR150993 sequence AR152416 sequence	AX026717 Sequence M74305 Synthetic e	A78762 Sequence 3 AR014735 Sequence	6 Seq	X04879 CaMV promot AJ308514 Synthetic	ğ i	AX405116 Sequence BD001990 A transge	AX033493 Sequence AX044092 Sequence	AF434748 Zea mays AF434750 Zea mays	AF434747 Zea mays AF434746 Zea mays	AF434749 Zea mays AX164073 Sequence	AX093047 Sequence	AX036754 Sequence	AX036755 Sequence AX036753 Sequence	AX036603 Sequence AX036739 Sequence	AX036616 Sequence	AX036611 Sequence	AX036758 Sequence	AX036759 Sequence	AX036741 Sequence	AX036601 Sequence	36610	Description

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RESULT 3
AX036601
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Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus

Patent: WO 0058485-A 13 05-OCT-2000;

MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR);

GRUBER VERONIQUE (FR)

Location/Qualifiers
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                                 synthetic construct.
synthetic construct
artificial sequences
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Sequence 7 from Patent FR2791358
AX036601
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                                                                                 AX036601.1
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  Patent:
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(bases 1 to 280)
nce, I., Theisen, M. and Gruber, V.
tent: FR 2791358-A 7 29-SEP-2000;
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                                                                                                                                                                                                                                                                                                                    /organism-"synthetic construct"
/db_xref-"texon:33630"
/note-"Directional desoxynucleotide building
/note-"14 c 17 g 16 t
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                                  sequences.
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59; Conservation
                                                                                                                                     398 bp
Sequence 6 from Patent W00058485
AX036740
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Sequence 7 from Patent W00058485
AX036741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           artificial sequences.

1 (bases 1 to 301)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating a
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 7 05-OCT-2000;
Rance, I., Theisen, M. and Gruber, V. Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 6 05-OCT-2000;
                                                    artificial sequences.
1 (bases 1 to 398)
                                                                                  synthetic construct.
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ilarity 100.0%;
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1 73 c 42 g
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                                                                                                                                                                                                                                                                                                                                                                                           /organism="synthetic construct"
/db_xref="taxon:32630"
/note="promoter MPr1154"
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/db_xref="taxon:32630"
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Pred. No. 3.9e-12;
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Pred. No. 3.8e-12;
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AX036758
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SOURCE
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AX036759
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Matches 59
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Best Local S
Matches 59
                                                          TITLE
                                                                                                          ORGANISM
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                                                                                                                                                                                                                                       Local Similarity
            artificial sequences.

1 (bases 1 to 541)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 24 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR)
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric expression promoters originating mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 25 05-OCT-2000;
                                                                                                                                                       AX036758
Sequence 24 from Patent AX036758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              artificial sequences.
1 (bases 1 to 472)
Rance, I., Theisen, M. and Gruber, V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
AX036759
                                                                                                        synthetic construct
                                                                                                                   synthetic construct.
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                                                                                                                                           AX036758.1 GI:11226267
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          GRUBER VERONIQUE (FR)
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ilarity 100.0%;
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ilarity 100.0%;
Conservative (
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/db_xref="taxon:32630"
/note="promoter MPr1169"
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/db_xref="taxon:32630"
/note="promoter MPr1147"
Location/Qualifiers
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WO0058485
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o. 3.8e-12;
o. 3.90 0;
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3.8e-12;
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                                                                                                                                                                              PAT 16-NOV-2000
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ACCESSION
VERSION
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AX036611
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VERSION
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Best Local S
Matches 59
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating imottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 23 05-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
AX036757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      synthetic construct. synthetic construct
                                                                                 artificial sequences.

1 (bases 1 to 79)

Rance,I., Theisen,M. and Gruber,V.

Patent: FR 2791358-A 17 29-SEP-2000;
                                                                                                                                synthetic construct
                                                                                                                                                                              Sequence 17
AX036611
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                                                          MERISTEM THERAPEUTICS (FR)
Location/Qualifiers
                                                                                                                                            synthetic construct.
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/organism="synthetic construct"
/db_xref="taxon:32630"
/note="Directional building block S7-Directional building block oligonucleotide for the construction of promoters by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /organism="synthetic construct"
/db_xref="taxon:32630"
/note="promoter MPr1168"
1...541
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                                                                                                                                                                    GI:11226206
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Pred. No.
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Pred. No.
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AX036616
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AUTHORS
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KEYWORDS
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AX036748
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ORIGIN
BASE COUNT
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Matches 46
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                                                                                           promoter
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Sequence 14 from Patent W00058485.
AX036748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct
artificial sequences.

1 (bases 1 to 79)
1 (bases 1 to 79)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating importie virus and cassava vein mosaic virus
Patent: WO 0058485-A 14 05-OCT-2000;
                                                                                                                                                                                                                                                                                Ax036616
Sequence 22 from Patent
Ax036616
                                                                                                                                                                                                                                                                                                                                                                                                                               46; Conservative 0;
                                                                                                                                                                     1 (bases 1 to 259)
Rance, I., Theisen, M. and Gruber, V.
Patent: FR 2791358-A 22 29-SEP-2000;
                                                                                                                                                                                                              synthetic construct artificial sequences.
                                                                                                                                                                                                                                         synthetic construct.
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                                                                                                                                            THERAPEUTICS (FR)
Location/Qualifiers
         /note="Promoter MPr1112 differs from MPr1111 by a deletion of 35 bp containing the 'G' box and stretching from position -127 to position -89 and a deletion of two bp situated at positions -78 and -76 promoter MPr1112"

a 67 c 38 g 72 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="synthetic construct"
/db_xref="taxon:32650"
/note="pirectional desoxynucleotide building
15 c 20 g 22 t
                                                                                                  /organism="synthetic construct"
/db_xref="taxon:32630"
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b; Pred. No. 5.1
0; Mismatches
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Pred. No.
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J. 5.1e-07;
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5.1e-07;
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DEFINITION
ACCESSION
VERSION
                                                                                                                                                                                                                                                                                                                      RESULT 13
AX036739
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SOURCE
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Sequence 9
AX036603
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46; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           artificial sequences.

1 (bases 1 to 296)
Rance,I., Theisen,M. and Gruber,V.
Patent: FR 2791358-A 9 29-SEP-2000;
MERISTEM THERAPEUTICS (FR)
                                                                                                                                                          Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 5 05-OCT-2000;
                                                                                                                                   MERISTEM THERAPEUTICS (FR); RANCE; GRUBER VERONIQUE (FR)
                                                                                                                                                                                                            artificial sequences.
1 (bases 1 to 371)
                                                                                                                                                                                                                                    synthetic construct
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AX036739
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llarity 100.0%;
Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note-"Promoter MPr1111 created by inserting at -99 bp position of MPr1098, an 18 bp element containing a 'G' and fusing a sequence of 58 bp (duplication of the elemes and as1)
                                                                   /organism="synthetic construct"
/db_xref="taxon:32630"
|note="promoter MPr1146"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               promoter MPr1111"
74 c 45 ç
                                                                                                                     Location/Qualifiers
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/db_xref="taxon:32630"
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                                                                                                                                                                                                                                                                                              371 bp
from Patent WO0058485
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 Score 46;
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4.7e-07;
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AX036755/c
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AX036753
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               Query Match
Best Local Similarity
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1 (bases 1 to 393)

Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus

Patent: WO 0058485-A 19 05-CCT-2000.

Patent: WO 0058485-A 19 05-CCT-2000.
                                                                                                                                                                                                                                                                                                                       393 bp
Sequence 19 from Patent WO0058485.
AX036753
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          artificial sequences.

1 (bases 1 to 392)

Rance,I., Theisen,M. and Gruber,V.

Chimeric expression promoters originating from commelina yellow
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 21 05-OCT-2000;
MERISTEM.THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR)
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Sequence 21 from Patent WO0058485.
AX036755
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             synthetic construct. synthetic construct
                                                                                                                                              MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR); GRUBER VERONIQUE (FR)
LOCATION/Qualifiers
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                                                                                                                                                                                                                                                                      synthetic construct
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71.4%; Score 45; DB 6; ilarity 100.0%; Pred. No. 1.2e-0 Conservative 0; Mismatches
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                                                                                      /organism="synthetic construct"
/db_xref="taxon:32630"
/note="promoter MPr1162"
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/db_xref="taxon:32630"
/note="promoter MPr1164"
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hes 0;
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Search completed: May 24, 2003, 14:52:24
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63
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/SIDS2/gcgdata/geneseq/geneseqn-embl/NA2001B.DAT:*
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AAA96841
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ALIGNMENTS

RESULT 1 AAA96847 ID AAA9

AAA96847 standard; DNA; 63 BP

AAA96847;

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Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; ss.
                 WPI; 2000-647238/62.
                                                                        29-MAR-1999;
                                                                                          29-MAR-2000; 2000WO-IB00370
                                                                                                             05-OCT-2000
                                                                                                                               WO200058485-A1.
                                                                                                                                                Synthetic.
                                                                                                                                                                                                               Directional desoxynucleotide building block
                                                                                                                                                                                                                                    19-FEB-2001
                                   Rance I,
                                                     (MERI-) MERISTEM THERAPEUTICS.
                                  Gruber V,
                                                                                                                                                                                                                                 (first entry)
                                                                        99FR-0003925.
                                   Theisen
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Chimeric expression promoter for transgenic plant production, comprises

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RESULT 2
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Matches 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       block, which was used to construct chimeric promoters of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                    Chimeric expression promoter for producing dicotyledonous an monocotyledonous transgenic plants comprises a nucleic acid derived from a promoter of the pea plastocyanin gene -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The
The present sequence represents the directional building block St
which is used to construct chimeric promoters of the invention in
PCR reactions. The specification describes a chimeric expression
promoter comprising a petE promoter of the pea plastocyanin gene.
                                                              Claim
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                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                     plastocyanin
                                                                                                                                                                                                                                                                                                                                                  petE promoter;
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                                                                                                                                                                                                                                                                                                                                               chimeric promoter; transgenic plant; MPr1108
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                                                                                                                                                                                                                                                                                                                                    promoter;
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RESULT 3
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                                                      The present sequence represents the chimeric promoter Mpr1110. The promoter is derived from the petE promoter from pea plastocyanin gene, by fusing the petE as-1 like and nos enhancer like elements to the promoter Mpr1098 (comprising TATA and CAAT boxes of petE), and then inserting a G box and fusing a fragmentof 35S RNA of CMV. The petE promoter directs cell-specific but not full light-regulated expression i transgenic tobacco plants. The promoter is used to construct chimeric expression promoters of the invention. The specification describes a chimeric comprising a G box operably or functionally linked upstream of a CAAT box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      comprising a G box operably or functionally linked upstream of a CAAT box. TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants; such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, rape, beetroot, or sunflower, and monocotyledonous species, e.g. wheat; barley, oat, rice, or corn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Chimeric expression promoter for producing dicotyledonous an monocotyledonous transgenic plants comprises a nucleic acid derived from a promoter of the pea plastocyanin gene .
                            dicotyledonous species, e.g. potato, tobacco, cotton, melon, cucumber, pea, rape, beetroot, or sunflower, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim
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Pred. No.
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                                                 Matches
                                                                                                                               The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                              with
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus transgenic plant; chimera; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide sequence of chimeric expression
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 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 CTGCAGACTAGTGGATTGATGTGATATCTCCCACTGACGTAAGGGATGACGCATGCCACT 63
            5 CTGCAGACTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 63
                                                                                                                      producing transgenic plants.
                                                                                                                                                                                                                                                                                                             ence from
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 CTGCAGACTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT
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                                                                                                                                                                                                                                                                                                          expression promoter for transgenic plant production, compris
from promoter comprising vascular expression region replaced
mence from promoter comprising green tissue expression region
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                                                                                                                                                                                                                                                                                                                                                                                     Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cassava
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larity 100.0%;
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                                                                                               A; 54 C;
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                                                                                               75 T; 0 other;
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1.4e-13;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chimeric - Commelina yellow mottle virus Chimeric - Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
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                                                                                                                                   AAA96859
                                                                                                                                                                                      AAA96859 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a chimeric promoter of the The specification describes chimeric expression promoters.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                        DNA;
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1.5e-13;
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Nucleotide sequence

of chimeric entry)

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Chimeric -
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                                                    Chimeric -
Chimeric -
                                                                                                      chimeric expression promoter; plant vascular expression plant green tissue expression promoter; Cassava vein mos
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                                                                                                                                                               Nucleotide sequence of chimeric expression promoter MPr1168
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                                                                                              transgenic
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                                                                                                                                    intergenic region; Commelina yellow
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                                                                                              plant;
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100.0%; Pr
                                                                                                                                                                                         entry)
                                                                   yellow mottle virus
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Best Local S
Matches 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                              WPI; 2000-647238/62
                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric - Commelia
Chimeric - Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA96857 standard; DNA; 604
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 5;
                                                                                                                                                                                                                                                 29-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nucleotide sequence of chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19-FEB-2001
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                                                                                    Rance I,
                                                                                                                                      (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                           29-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            transgenic plant;
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                                                                                    Gruber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gruber
                                                                                                                                                                                                                                                                                                                                                                                                                                       Commelina yellow mottle virus
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                                                                                                                                                                                                                                                 2000WO-IB00370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               chimera;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    91pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                mosaic virus
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Z
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             promoter MPr1167
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Chimeric expre sequence from with sequence

expression promoter for transgenic plant production, comprises from promoter comprising vascular expression region replaced uence from promoter comprising green tissue expression region

The Claim

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Page 87;

91pp; English

The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful

sequence represents a chimeric promoter of the invention cation describes chimeric expression promoters. These

for producing transgenic plants.

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RESULT 9
AAA96848
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Best Local S
Matches 59
                                                                                 Chimeric expression promoter for transgenic plant production, compris sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                    The
  from a first
                                      The
                                                        Disclosure;
                                                                                                                          WPI;
                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                                                               chimeric expression promoter;
                                                                                                                                                                                                                                                                                                                            Directional
                            block,
                                                                                                                                             Rance
                                                                                                                                                               (MERI-)
                                                                                                                                                                                                    29-MAR-2000;
                                                                                                                                                                                                                      05-OCT-2000
                                                                                                                                                                                                                                                                                      plant green tissue expression
                                                                                                                                                                                                                                                                                                                                             19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                 AAA96848;
                                                                                                                                                                                                                                                                                                                                                                                  AAA96848 standard;
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                                                                                                                                                                                 29-MAR-1999;
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                                                                                                                                                                                                                                                                             transgenic plant;
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present sequence represents a directional desoxynucleotide building tok, which was used to construct chimeric promoters of the invention. specification describes chimeric expression promoters. These meric promoters comprise a nucleic acid sequence which is derived m a first plant promoter, in which a plant vascular expression
                                                                                                                                                                                                                                                                                                                                                                                                                                 CTGCAGACTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 63
                                                                                                                                                               MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          604 BP; 186 A; 116 C; 145 G;
                                                                                                                                                                                                                                                                                    intergenic region; Commelina yellow mottle virus; expression promoter; plant vascular expression proentissue expression promoter; Cassava vein mosaic
                                                                                                                                           Gruber V,
                                                        Page
                                                                                                                                                                                                                                                                                                                          desoxynucleotide building block
                                                                                                                                                                                                    2000WO-IB00370
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                                                        26;
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100.0%;
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GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT

Query Match Best Local S Matches 46

Similarity

Conservative

73.0%; Sur 100.0%; Pr 0;

Score 46; Pred. No.

DB 21;

Length 79;

1.3e-08;

Mismatches

Indels

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RESULT 10
ANA96472
ID ANA96472
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                                                     The present sequence represents the directional building block $7, which is used to construct chimeric promoters of the invention in PCR reactions. The specification describes a chimeric expression promoter comprising a petE promoter of the pea plastocyanin gene, o comprising a G box operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotto lettuce, tomato, melon, cucumber, pea, rape, beetroot, or sunflower and monocotyledonous species, e.g. wheat, barley, oat, rice, or correct the production of the pea plants, such as dicotyledonous species, e.g. wheat, barley, oat, rice, or correct the pea plants, such as dicotyledonous species, e.g. wheat, barley, oat, rice, or correct the pea plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava veln mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                             Chimeric expression promoter for producing dicotyledonous monocotyledonous transgenic plants comprises a nucleic aci derived from a promoter of the pea plastocyanin gene
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     Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleotide sequence of the directional building block S7.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         promoter; chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2000-587667/55
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                                                                                                                                                                                                                                                                                                                                                                        24; Page 76; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        46; Conserv
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     B₽;
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  22 A; 15 C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        promoter;
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  20 G;
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Pred. No.
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     22 T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              79
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                63
                                                                                        sunflower,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
                                                                                                                   cotton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                      or
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RESULT 11
ANA96477
ID ANA96477
AC ANA96477
AC ANA96477
AC ANA96
AC ANA96477
ANA964477
ANA96447
ANA9647
ANA964
                                                                            AAA96465
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   ZXXX
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                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                     Query Match
Best Local (
                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents the chimeric promoter Mprill2. The promoter is derived from the petE promoter from pea plastocyanin gene, by fusing the petE as-1 like and nos enhancer like elements to the promoter Mpril098 (comprising TATA and CAAT boxes of petE), and then fusing a fragment comprising a duplication of the element as2 and as1. The petE promoter directs cell-specific but not full light-regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a Gox operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors
   08-FEB-2001
                                         AAA96465;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chimeric expression promoter for producing dicotyledonous and monocotyledonous transgenic plants comprises a nucleic acid so derived from a promoter of the pea plastocyanin gene .
                                                                            AAA96465 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 25; Page 79; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-587667/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               22-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-SEP-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO200056906-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pisum sativum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     plastocyanin gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     petE promoter; chimeric promoter; transgenic plant; MPr1112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nucleotide sequence of the promoter MPr1112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAA96477;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAA96477 standard; DNA; 259
                                                                                                                                                                                                                                                                                                                                                                                                  for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, p
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                          42
                                                                                                                                                                                                            18 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     34
                                                                                                                                                                    GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 79
                                                                                                                                                                                                                                                                                                                                                                                   beetroot,
                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                              259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gruber V,
                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                          BP; 82 A; 67 C;
                                                                                                                                                                                                                                                                                                                                                                 rice, or corn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2000WO-IB00317
 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99FR-0003635
                                                                                                                                                                                                                                                                                                                                                                                 or sunflower,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     promoter;
                                                                              DNA;
                                                                                                                                                                                                                                                                 73.0%;
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                                                                              296
                                                                            ВP
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                                                                                                                                                                                                                                                                   Score 46;
Pred. No.
                                                                                                                                                                                                                                                                                                                          38 G;
                                                                                                                                                                                                                                                                                                                                                                               and monocotyledonous
                                                                                                                                                                                                                                                 Mismatches
                                                                                                                                                                                                                                                                                                                          72 T;
                                                                                                                                                                                                                                                                 DB 21; ; 1.9e-08;
                                                                                                                                                                                                                                                                                                                            0
                                                                                                                                                                                                                                                                                                                            other;
                                                                                                                                                                                                                                                                                    Length 259;
                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                   species,
                                                                                                                                                                                                                                                                                                                                                                                                  cucumber, pea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence
                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                 RESULT 13
AAA96839
OS XXX XXX DE XXX OS XXX
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
                                                                                                                                                                                                                                                                                                                                                                               S
                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local 9
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 Chimeric
                                     transgenic plant;
                                                                                                                                 Nucleotide
                                                                                                                                                                      19-FEB-2001
                                                                                                                                                                                                          AAA96839;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 296 BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-587667/55
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200056906-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          petE promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    barley, oat,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       e.g. potato, to
rape, beetroot,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pisum sativum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          28-SEP-2000
                                                                                                                                                                                                                                                                                                                                             42
                                                                                                                                                                                                                                                                                                                                                                 18 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT
                                                                                                                                                                                                                                                                                                                                             GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT
                                                                                                                                                                                                                                                                                                                                                                                                                      46;
                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gruber
                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99FR-0003635
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94 A; 74 C;
                                     chimera;
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20-MAR-2000; 2000WO-IB00317
                                                                                                                                                                    plastocyanin gene promoter;
                                                                                                                                                                                                                         Nucleotide sequence of the
                                                                                                                                                                                  chimeric promoter;
                                                                                                                                                                                                                         promoter
                                                                                                                                                                    88
                                                                                                                                                                                    transgenic plant; MPrllll;
                                                                                                                                                                                                                         MPr1111
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Chimeric expression promoter for producing dicotyledonous ar monocotyledonous transgenic plants comprises a nucleic acid derived from a promoter of the pea plastocyanin gene sequence

Claim 2; Page 71; 83pp; English.

full light_regulated expression in transgenic tobacco plants. The promoter is used to construct chimeric promoters of the invention. The specification describes a chimeric expression promoter comprising a promoter of the pea plastocyanin gene, or comprising a G box operably or functionally linked upstream of a CAAT box, TATA box and transcription initiation site. The chimeric promoters are used in expression vectors for producing transgenic plants, such as dicotyledonous species, e.g. potato, tobacco, cotton, lettuce, tomato, melon, cucumber, pea, promoter is derived from the petE promoter from pea plastocyanin gene, by fusing the petE as-1 like and nos enhancer like elements to the promoter MerI1098 (comprising TATA and CAAT boxes of petE), and then inserting a G box and fusing a fragment comprising a duplication of the element as2 and as1. The petE promoter directs cell-specific but not full that the petE promoter directs cell-specific but not rice, or corn. or sunflower, represents the and monocotyledonous promoter Mpr1111. species, wheat,

45 G; 83 T; 0 other;

73.0%; 0, Score 46; Pred. No. Mismatches DB 21; 2e-08; 0 Length 296; Indels 63 0 Gaps 0

87

AAA96839 standard; DNA; 371 BP.

sequence of chimeric expression promoter MPr1146.

Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus virus;

Commelina yellow mottle virus

05-OCT-2000 WO200058485-A1 Chimeric -

2000WO-IB00370

Cassava

vein mosaic virus.

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AAA96855/C
ID AAA96855/C
ID AAA9685/C
XX AA96
XX IP-FE
XX Nucle
XX Promc
KW Promc
KW Promc
KW Plant
KW Plant
KW Plant
KW Plant
XX OS-OC
XX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI;
                                                                                                                                                    05-OCT-2000
                                                                                                                                                                                                                                                           Chimeric - Chimeric -
                                                                                                                                                                                                                                                                                                                                                                Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA96855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  With
                                                      29-MAR-1999;
                                                                                                    29-MAR-2000;
                                                                                                                                                                                                           WO200058485-A1
                                                                                                                                                                                                                                                                                                                                       transgenic plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleotide sequence of chimeric expression promoter MPr1164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAA96855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       eric expression promoter for transgenic plant production, comprise
ence from promoter comprising vascular expression region replaced
sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2000-647238/62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               . Similarity
46; Conserv
  MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; DNA; 392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page 81;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gruber V,
                                                                                                                                                                                                                                                      Commelina yellow mottle virus Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                             tissue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            73.0%; Solarity 100.0%; I Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP;
                                                                                                    2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99FR-0003925.
                                                99FR-0003925.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 122 A; 68 C;
                                                                                                                                                                                                                                                                                                                                       chimera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Theisen
                                                                                                                                                                                                                                                                                                                                          88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 46;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                             plant vasc
promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 89 G; 92 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 21; I
2.1e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 371;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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sequence with sequ

Claim 5;

Page

85;

91pp;

English

Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    sequence
with sequ
                                                                                                                                                                                                                                                                           chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                           Chimeric -
                                                                                                                                                                                                                                                                                                                                   Nucleotide sequence of chimeric expression promoter MPr1162.
                                                                                                                                                                                                                                                                                                                                                                                  AAA96853;
                                                                                                                                                                                                                                                                                                                                                                                                      AAA96853 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric
                                                                                                                                                   29-MAR-1999;
                                                                                                                                                                        29-MAR-2000;
                                                                                                                                                                                               05-OCT-2000.
                                                                                                                                                                                                                    WO200058485-A1
                                                                                                                                                                                                                                                      Chimeric -
                                                                                                                                                                                                                                                                                                               Promoter;
                                                                                                                                                                                                                                                                                                                                                           19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
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                                                                                                                           (MERI-) MERISTEM THERAPEUTICS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page
                                                                                                   Gruber V,
                                                                                                                                                                                                                                                                                                             intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gruber V,
                                                                                                                                                                                                                                          Cassava v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        71.4%; Sillarity 100.0%; Conservative 0;
                                                                                                                                                                        2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BP; 127
                                                                                                                                                                                                                                                                                                                                                          (first
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             transgenic plants.
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                                                                                                     Theisen
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                                                                                                                                                                                                                                           mosaic
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0; Mismatches
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x333333333333x**x x** The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.

Sequence 393 BP; 128 A; 75 C; 93 G; 97 T; 0 other;

ş Query Match 71.4%; Score 45; DB Best Local Similarity 100.0%; Pred. No. 5.3 Matches 45; Conservative 0; Mismatches Score 45; DB 21; Length C., pred. No. 5.3e-08; indels Length 393; Gaps

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Search completed: May 24, 2003, 14:34:44 Job time: 109.726 secs

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Result
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Maximum DB
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1: /cgn2_6/ptodata/1,
2: /cgn2_6/ptodata/1,
3: /cgn2_6/ptodata/1,
4: /cgn2_6/ptodata/1,
5: /cgn2_6/ptodata/1,
6: /cgn2_6/ptodata/1,
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Match
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63
  Listing first 45 summaries
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/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/BCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/backfiles1.seq:*
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                                                     US-09-329-169-1

US-09-330-714A-1

US-09-328-826-1

US-09-042-426-5

US-09-291-238-5

US-09-330-760-5

US-09-330-769-5

US-09-330-714A-5

US-09-330-714A-5

US-09-330-714A-5

US-09-328-826-5

US-09-328-826-5

US-09-328-826-5

US-09-328-7736-1

US-08-897-736-1
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US-09-291-238-1

US-09-330-760-1

US-09-328-473-1

US-09-330-737-1
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-08-764-100-23
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 8-446-486-31
8-463-308-31
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US-08-247-809A-3
                                                       US-08-247-809A-3
                      Query Match
Best Local
             Matches
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                                                                  TOPOLOGY: 1
MOLECULE TYPE:
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                                                                                                            LENGTH:
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43.4			43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4
68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9
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US-08-525-507-16	US-07-966-187-17	US-08-711-728-5	US-08-247-809A-5	US-09-065-999-6	US-09-065-999-5	US-09-458-093-2	US-08-894-440-2	US-09-471-913-6 ·	US-09-185-244-2	US-08-729-601A-46	US-09-011-151-9	US-09-011-151-8	US-09-363-970-35	US-08-729-601A-43	US-07-936-163-46	5254799-30	US-08-463-308-30
Sequence 16	Sequence 17	Sequence 5,	Sequence 5,	Sequence 6,	Sequence 5,	Sequence 2,	Sequence 2,	Sequence 6,	Sequence 2,	Sequence 46	Sequence 9,	Sequence 8,	Sequence 35	Sequence 43	Sequence 46,	Patent No. 5	Sequence 30,
5, App	7, App	, Appl	, Appl	, Appl:	, Appli	, Appli	, Appli	-	-	5, Appi	, Appli	⊳	5, Appl		5, Appl	in), Appi

ALIGNMENTS

Sequence 3, Application US/08247809A Patent No. 5569823 SOFTWARE: WordPerfect 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/247,809A FILING DATE: May 23, 1994 CLASSIFICATION: 800 PRIOR APPLICATION DATA: APPLICATION NUMBER: P 43 178 45.6 (G COUNTRY: U.S.A. ZIP: 10591-5144 COMPUTER READABLE FORM: MEDIUM TYPE: blskette, MEDIUM TYPB: storage INFORMATION FOR SEQ ID NO: REGISTRATION NUMBER: 33,141 REFERENCE/DOCKET NUMBER: Bayer TELECOMMUNICATION INFORMATION: TELEPHONE: (914) 332-1700 TITLE OF INVENTION: DEOXYRIBONUCLEIC ACIDS NUMBER OF SEQUENCES: 18 CORRESPONDENCE ADDRESS: APPLICATION NUMBER: P 43 FILING DATE: May 28, 199 ATTORNEY, AGENT INFORMATION: NAME: Kurt G. Briscoe REGISTRATION NUMBER: 33, SEQUENCE CHARACTERISTICS: APPLICANT: Peter H. Sci APPLICANT: Edgar Maiss COMPUTER: NEC Powermat OPERATING SYSTEM: DOS TELEFAX: (914) 332-1844 nucleic acid E: SPRUNG HORN KRAMER & WOODS 660 White Plains Road Diskette, 3.50 inch, DNA (genomic) single Powermate 1 Plus 68.9%; 97.8%; Schreier; Klaus Stenzel; ω •• Score 43.4; Pred. No. 4 45.6 (Germany) 9049-KGB 2.0 ¥ DB Gunter Adam; Length 439;

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Mismatches

Indels

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                                                                Patent No. 5773700
GENERAL INFORMATION:
                                                                                             Sequence 23,
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FILING DATE: 23-MAY-1
PRIOR APPLICATION DATA:
APPLICATION NUMBER: D
FILING DATE: 28-MAY-1
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APPLICANT:
                                     APPLICANT:
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           APPLICANT:
                                                                                                                                                              FITLE OF INVENTION: DEOXYRIBONUCLEIC ACIDS NUMBER OF SEQUENCES: 18
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                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
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REFERENCE/DOCKET NUMBER: Bayer 9049.1-KGB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/711,728 FILING DATE: 03-SEPT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM:
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10591-5144
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                                                                                                                                                                                                                                                                                                                              nucleic acid
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                                                                                            Application US/08764100
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                                                                                                                                                                                                                                                                                                                                             439 base pairs
          De Haan, Petrus T.
Gielen L., Johannes J.
Peters, Dirk
                                                                                                                                                                                                                                                                                                                                                                                                    (914) 332-1844
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                                                  van Grinsven J., Martinus
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97.8%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.50 inch,
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Pred. No. 4.3e-09;
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RESULT 4
US-09-042-426-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local :
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FILING DATE: 17-MAR-1993
APPLICATION NUMBER: GB 9206016.9
FILING DATE: 19-MAR-1992
ATTORNEY/AGENT INFORMATION:
NAME: NO. 5773700115, Allen E.
REGISTRATION NUMBER: 34,490
REFERENCE/DOCKET NUMBER: 137-1061
TELECOMMUNICATION INFORMATION:
THELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (415) 857-11
INFORMATION FOR SEQ ID NO:
                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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MEDIUM TYPE: Floppy disk
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NUMBER OF SEQUENCES:
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les 44; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                            New Jersey
                                                                                                                                                                                                             E: No. 6114608artis Corporation 564 Morris Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 446 base pairs
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                                                                                                                                                                                                                                                                                                 Irvin J.
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MBER: US/09/042,426
March 13, 1998
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                                                                                                                                                                                                                                                                                Mettler, Paul
DNA Construct
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Pred. No. 4.
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                                                                                                                                                                                                                                                                                S. Dietrich, Ralph Sinibaldi
Containing Bacillus
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ATTORNEY/AGENT INFORMATION:

135/1

CLASSIFICATION:

800

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RESULT 5
US-09-291-238-1
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                                                            TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Description US/09291238 Patent No. 6222104
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Best Local :
                                                                                                                                                            APPLICATION NUMBER: 09/042,42
FILING DATE: MATCh 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,93
REFERENCE/DOCKET NUMBER: 135/
TELECOMMUNICATION INFORMATION:
TELECPHONE: (919) 541-8614
TELECPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Irvin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
WENTUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6222104artis Corporation
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REGISTRATION NUMBER: 32,993
REFERENCE, DOCKET NUMBER: 13
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION:
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CLONE: 35S Pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: D. HYPOTHETICAL: NO
                            MOLECULE TYPE:
               HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Summit STATE: New T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    268 CAAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 312
                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 CTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 532 base pairs TYPE: nucleic acid STRANDEDNESS: double TOPOLOGY: linear
                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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VENTION: DNA Construct Containing Bacillus
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                              DNA (genomic)
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l3, 1998
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Pred. No. 4.6e-09;
0; Mismatches 1;
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RESULT 7
US-09-328-473-1
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US-09-330-760-1
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                                                                                                                              Matches
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CLONE: 35S Promoter
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APPLICATION NUMBER: 09/0,
FILING DATE: MArch 13, 1:
ATTORNEY/AGENT INFORMATION:
NAME: Hoxie, Thomas
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6229075art1s Corporation
                                                                                                                                                                                                                                        MOLECULE TYPE: DI
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
                                                                                                                                                                                                          IMMEDIATE SOURCE:
CLONE: 35S Promoter
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TELECOMMUNICATION INFORMATION:
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TITLE OF INVENTION:
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STATE: New Jersey
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                                                                                                                              Local Similarity 97.8 
les 44; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Hoxie, Thomas REGISTRATION NUMBER: 32,993
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/ENTION: DNA Construct Containing Bacillus
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                                                                                                                                          Score 43.4; DB 4;
Pred. No. 4.6e-09;
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Pred. No. 4.6e-09;
                                                                                                                              Mismatches
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                                                                                                                                                              DB 4;
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Sequence 1, Application US/09328473 Patent No. 6232533

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us-09-330-737-1
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                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
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                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                        NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6232534art1s Corporation
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HYPOTHETICAL: P
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
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APPLICATION NUMBER: 0:
FILING DATE: March 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STREET: SUMMIT CITY: Summit CITY: New Jersey
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ADDRESSEE: No. 6232533artis Corporation
STREET: 564 Morris Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: Hox1e, Thomas
REGISTRATION NUMBER: 32,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                              APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IMMEDIATE SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                         ZIP: 0790
                                                                                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                         12 CTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/09/328,473 FILING DATE:
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                                                                                                      Summit
: Summit
: New Jersey
:RY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : 532 base pairs nucleic acid
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                                                                                                                                                          564 Morris Avenue
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PatentIn Release #1.0, Version #1.25
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ENTION: DNA Construct Containing Bacillus
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Pred. No. 4.6e-09;
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US-09-329-169-1
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GENERAL INFORMATION:
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Best Local
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INFORMATION FOR SEQ ID NO:
                                   INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 532 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
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CLONE: 35S Promoter
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: March 13, 1 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     268 CAAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12 CTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 32,993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:
                                                                    NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
                                                                                                                                                                                                                                            PRIOR APPLICATION DATA
                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/329,169
FILING DATE: 09-Jun-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                       APPLICATION NUMBER: 09/042,426 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Application US/09329169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Summit
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
             ENGTH: 532 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      564 Morris Avenue
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Pred. No. 4.6e-09
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US-09-330-714A-1
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                                                                                                                                                  Query Match
                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (919) 541-8
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Irv1
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                                                                                                                Local Similarity
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12 CTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
                                                                                                                                                                                                                  ANTI-SENSE: ...
IMMEDIATE SOURCE:
CLONE: 35S Promoter
CLONE: 35S P
                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: HOXIE, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/330,714A
FILING DATE: 11-Jun-1999
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 634
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CLONE: 35S Promoter
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
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TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 09/042,426 FILING DATE: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 532 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                              Conservative
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NVENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             564 Morris
                                                                                                         68.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6342660artis Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             541-8689
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                                                                                                         Score 43.4; DB 4;
Pred. No. 4.6e-09;
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                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
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                                                                                                                                                  DB 4;
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                                                                                                                                              Length 532;
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                                                                                                                                                                                            RESULT 12
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US-09-328-826-1
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                                                                                                                                   Sequence 5, Application US/09042426 Patent No. 6114608
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                                                                                                                                                                                                                                                                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                  APPLICANT: Irvin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
               NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6114608artis Corporation
                                                                            TITLE OF INVENTION:
                                                                                                                                                                                                                                               268 CAAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 312
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12 CTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 56
 STREET:
                                                                                                                                                                                                                                                                                                                          Local Similarity les 44; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/328,826
EILING DATE: 09-Jun-1999
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
                                                                                                                                                                                                                                                                                                                                                                                                                  CLONE: 35S Promoter SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS
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HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Hoxie, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/042,426 FILING DATE: March 13, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: New Jersey
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 532 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DDRESSEE: NO.
564 Morris Avenue
                                                                            Irvin J. Mettler, Paul S. Dietrich, Ralph Sinibaldi
VENTION: DNA Construct Containing Bacillus
                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    564 Morris Avenue
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                                                                                                                                                                                                                                                                                                                                       Pred.
                                                                                                                                                                                                                                                                                                                                           Score 43.4;
Pred. No. 4
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                                                                                                                                                                                                                                                                                                                                                         DB 4;
                                                                                                                                                                                                                                                                                                                                                         Length 532;
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US-09-291-238-5
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5, Applic Patent No. 6222104
            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/042,426
FILING DATE: MARCH 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOXLE, Thomas
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                       COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: 35S Promoter
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TITLE OF INVENTION:

NUMBER OF SEQUENCES:
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RELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                           CLASSIFICATION:
                                                                                                                                                            FILING DATE:
                                                                                                                                                                         APPLICATION NUMBER: US/09/291,238
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bes 44; Conserv
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REGISTRATION NUMBER: 32,993
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DNA Construct Containing Bacillus
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Pred. No. 4.7e-09;
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Query Match
Best Local Similarity
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GENERAL INFORMATION:
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SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                        TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO: 5:
                                                                                       MOLECULE
HYPOTHETICAL: N
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APPLICATION NUMBER:
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ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: 35S Promoter
                                                                       ANTI-SENSE: NO
IMMEDIATE SOURCE:
                                                                                                                    MOLECULE TYPE:
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CORRESPONDENCE ADDRESS:
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TITLE OF INVENTION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Pelesco
                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                LENGTH: 560 base pairs
TYPE: nucleic acid
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CITY: Summit
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                                                                                                                                     TOPOLOGY:
                                                                                                                                                                                                                                                                                       REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 07901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE:
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/ENTION: DNA Construct Containing Bacillus
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                                                                                                                      DNA (genomic)
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541-8689
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97.8%;
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97.8%;
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                                                                                                                                                                                                                                                                                       32,993
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 Score 43.4; DB 4; Pred. No. 4.7e-09;
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               Length 560;
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MOLECULE TYPE: DNA (9en
HYPOTHETICAL: NO
ANTI-SENSE: NO
IMMEDIATE SOURCE:
CLONE: 35S Promoter
US-09-328-473-5
Search completed: May 24, 2003, 15:37:43 Job time: 28.0806 secs
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US-09-328-473-5
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                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/042,426
FILING DATE: March 13, 1998
ATTORNEY/AGENT INFORMATION:
NAME: HOX1e, Thomas
REGISTRATION NUMBER: 32,993
REGISTRATION NUMBER: 32,993
REFERENCE/DOCKET NUMBER: 135/1
TELECOMMUNICATION:
TELEPHONE: (919) 541-8614
TELEPHONE: (919) 541-8689
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 560 base pairs
TYPE: DULLeic acid
TYPE: DULLeic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: DNA (GENOMIC)
                                                                                                                                    Query Match 68.9%;
Best Local Similarity 97.8%;
Matches 44; Conservative
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APPLICANT: Irvin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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TITLE OF INVENTION: DNA Construct Containing Bacillus
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: No. 6232533artis Corporation
STREET: 564 Morris Avenue
                                                                STREET: 564 Morris
CITY: Summit
STATE: New Jersey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/328,473 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 07901
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                                                                                                                                    Score 43.4; DB 4;
Pred. No. 4.7e-09;
0; Mismatches 1;
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Minimum DB
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Perfect score:
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Maximum Match 100%
Listing first 45 summaries
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      43.4
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length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /cgn2_6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/USOB_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/USOB_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/USO9_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/US06_PUBCOMB.seq:*
/cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*
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/cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US10_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US60_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*
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                                                                392
392
393
393
462
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566
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                                       US-09-963-803-7
US-09-963-803-7
US-09-963-803-25
US-09-963-803-24
US-09-963-803-14
US-09-963-803-14
US-09-963-803-1
US-09-963-803-1
US-09-963-803-1
US-09-963-803-20
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1693.834 Million cell updates/sec
                                                                                  Sequence 22, Appl
Sequence 47, Appl
Sequence 3, Appli
Sequence 33, Appl
                                                                                                                       Sequence 13, Appl
Sequence 7, Appli
Sequence 6, Appli
Sequence 25, Appli
Sequence 24, Appl
Sequence 24, Appl
Sequence 14, Appl
Sequence 5, Appli
Sequence 21, Appl
Sequence 21, Appl
Sequence 20, Appl
Sequence 22, Appl
Sequence 22, Appl
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39	39	41.8	41.8	41.8	41.8	41.8	41.8	41.8	41.8	41.8	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4	43.4
61.9	61.9	66.3	66.3	66.3	66.3	66.3	66.3	66.3	66.3	66.3	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9	68.9
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US-09-870-375-28	US-09-870-375-33	US-10-055-001A-13	US-10-055-001A-23	US-10-055-001A-26	US-10-055-001A-24	US-10-055-001A-25	US-09-509-945-4	US-09-509-945-5	US-09-970-921-10	US-09-970-921-7	US-09-810-861B-4	US-10-074-279-10	US-09-316-622-7	US-09-316-622-6	US-09-316-622-8	US-09-316-622-8	US-09-316-622-5	US-09-316-622-5	US-10-047-542-101	US-09-316-622-1	US-09-316-622-2	US-09-316-622-4	US-09-316-622-3	US-09-810-861B-3	US-09-990-659A-15
28,	Sequence 33, Appl	Sequence 13, Appl	Sequence 23, Appl	Sequence 26, Appl	Sequence 24, Appl	Sequence 25, Appl	Sequence 4, Appli	Sequence 5, Appli	10,	Sequence 7, Appli	Sequence 4, Appli.	Sequence 10, Appl	Sequence 7, Appli	Sequence 6, Appl1	Sequence 8, Appli	Sequence 8, Appli	Sequence 5, Appli	Sequence 5, Appli	Sequence 101, App	Sequence 1, Appli	Sequence 2, Appli	Sequence 4, Appli	Sequence 3, Appli	Sequence 3, Appli	Sequence 15, Appl

ALIGNMENTS

US-09-963-803-13

Sequence 13, Application US/09963803 Publication No. US20030028922A1 GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: Virus and cassava vein mosaic virus

FILE REFERENCE: 184332042

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; OTHER INFORMATION: Directional desoxynucleotide building block ; OTHER INFORMATION: S6 US-09-963-803-13
RESULT 2
US-09-963-803-7
; Sequence 7, Application US/09963803
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                                                                                                                                                                                                                              Query Match 100.0%;
Best Local Similarity 100.0%;
Matches 63; Conservative 0
                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1 SEQ ID NO 13 LENGTH: 63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/963,803.
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                          61
                                                                                                                         61
                                                                                                                                                                                           1 CATGCTGCAGACTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC 60
                                                                                                                                                              CATGCTGCAGACTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCC
                                                                                                                         ACT 63
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Pred. No. 3e-16;
D; Mismatches
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                                                                                   Matches
                                                                                                                         Query Match
                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 39
SOFTWARE: PATENTIN VETSION 3.1
SEQ ID NO 6
LENGTH: 398
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                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: ER 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR PPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: PatentIn version 3.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: Chimeric expression promoters originating from commelina TITLE OF INVENTION: virus and cassava vein mosaic virus
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ORGANISM: Artificial Sequence
                                                                                                                                                                                   NAME/KEY: promoter LOCATION: (1)..(398) OTHER INFORMATION:
                                                                                                                                                                                                                                                FEATURE:
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                                                                                                                                                                                                                                                                                      FEATURE:
                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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OTHER INFORMATION: promoter MPr1154
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CTGCAGACTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 63
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                                                                                                      Similarity
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; Pred. No. 2.1e-14;
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Pred. No.
                                                                               Mismatches
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. 2.2e-14;
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; NAME/KEY: promoter
; LOCATION: (1)..(472)
; OTHER INFORMATION:
US-09-963-803-25
                                                                               US-09-963-803-24
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Publication No. US20030028922A1
GENERAL INFORMATION:
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APPLICANT: Meristem Therapuetics
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: virus and cassava vein mosaic virus
                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
SEQ ID NO 24
LENGTH: 541
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 59; Conserv
  Matches
                  Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: Chimeric expression promoters originating TITLE OF INVENTION: virus and cassava vein mosaic virus FILE REFERENCE: 184332042
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                                                                                                  NAME/KEY: promoter LOCATION: (1)..(54: OTHER INFORMATION:
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OTHER INFORMATION: promoter MPr1169
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                                                                                                                                                                               OTHER INFORMATION: promoter MPr1168
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Conservative
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                  93.7%;
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                  Score 59;
Pred. No.
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Pred. No. 2.4e-14;
Mismatches
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; FEATURE:
; NAME/KEY: promoter
; LOCATION: (1)..(604)
; OTHER INFORMATION:
US-09-963-803-23
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                                                                                                                                  ; OTHER INFORMATION: ; OTHER INFORMATION: US-09-963-803-14
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Sequence 14, Application US/09963803
Delication US20030028922A1
Delication US20030028922A1
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APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                              Query Match
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Publication No.
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                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEO ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
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CURRENT FILING DATE: 2001-09-26
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 39
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                   Q ID NO 14
LENGTH: 79
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                                                                                                                                                                                         FEATURE:
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                    18 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 63
                                                                            Local Similarity
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Similarity
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b. US20030028922A1
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                                                          Conservative
                                                                                                                                                        Directional desoxynucleotide building block S7
                                                                          73.0%;
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100.0%; Pred. No.
tive 0; Mismatc
                                                        0;
                                                                          Score 46; DB 9; 1; Pred. No. 2.8e-09;
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                                                          Mismatches
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                                                                                            Length 79
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                                                                                                                                                                                                                               SOFTWARE: Pate SEQ ID NO 21 LENGTH: 392 TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 21, Application US/09963803 Publication No. US20030028922A1
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Best Local :
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                                                                                                                              FEATURE:
NAME/KEY: promoter
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; FEATURE;
; NAME/KEY: promoter
; LOCATION: (1)..(371)
; OTHER INFORMATION:
US-09-963-803-5
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; Sequence 5, Application US/09963803
; Publication No. US20030028922A1
; GENERAL INFORMATION:
; APPLICANT: Meristem Therapuetics.
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                                                                                        Best Local Similarity 100.0%; | Matches 46; Conservative 0;
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CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
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                                                                                                                                                                                                                                                                                                                                                  LENGTH: 371
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                            FEATURE:
OTHER INFORMATION: promoter MPr1146
                          18 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 63
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93 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCACT 138
                                                                                     73.0%; Score 46; DB
100.0%; Pred. No. 4.5
ive 0; Mismatches
                                                                                                                DB 9; L
                                                                                                                                    Length 371;
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                                                                                          Gaps
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; LOCATION: (1)..(392); OTHER INFORMATION: US-09-963-803-21
Query Match 71.4%; Score 45; DB 9;
Best Local Similarity 100.0%; Pred. No. 1.2e-(
Matches 45; Conservative 0; Mismatches
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CURRENT FILING DATE: 2001-09-26
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT 1800/00370
PRIOR FILING DATE: 2000-10-05
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                                                                                                                                                                                                                                               OTHER INFORMATION: promoter MPr1164
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                                 1.2e-08;
                                                        Length 392;
        Indels
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Sequence 20, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
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SOFTWARE: PAtentin version 3.1
SEQ ID NO 19
LENGTH: 393
TYPE: DNA
ORGANISM: Artificial Sequence
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PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow TITLE OF INVENTION: virus and cassava vein mosaic virus FILE REFERENCE: 184332042 CURRENT APPLICATION NUMBER: US/09/963,803 CURRENT FILING DATE: 2001-09-26
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PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT 1800/00370
PRIOR FILING DATE: 2000-10-05
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CURRENT FILING DATE: 2001-09-26
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                                                                                                                                                 LENGTH: 462
TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                            FEATURE:
OTHER INFORMATION: promoter MPr1163
                                              OTHER INFORMATION:
                                                           NAME/KEY: promoter LOCATION: (1)..(462)
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                                                                                                        FEATURE:
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es 45; Conservative
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100.0%; Pr
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Pred. No.
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. 1.2e-08;
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US-10-012-070A-47
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                                                             ; TYPE: DNA
; ORGANISM: Cauliflower mosaic virus
US-10-012-070A-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 47, Application US/10012070A Publication No. US20030077801A1
                                                                                                                                              SEQ ID NO 47
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Best Local Similarity
                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PCT/GB00/01573
PRIOR FILING DATE: 2000-04-20
NUMBER OF SEQ ID NOS: 57
                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/012,070A
CURRENT FILING DATE: 2001-10-29
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                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Hawkes, Timothy APPLICANT: Warner, Simon
                                                                                                                                                                    SOFTWARE: PatentIn
                                                                                                                                                                                                                                                                                           FILE REFERENCE: 50490/UST
                                                                                                                                                                                                                                                                                                          APPLICANT: Pickerill, Andrew
TITLE OF INVENTION: Herbicide Resistant Plants
                                                                                                                         LENGTH: 309
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Warner, Simon Andrews, Christopher Bachoo, Satvinder

Ver.

2.0

68.9%; 97.8%;

Score 43.4; Pred. No. 4

4; DB 9; 4.9e-08;

Length 309;

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; OTHER INFORMATION: US-09-963-803-22
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                                                          Query Match
Best Local Similarity
Matches 45; Conserv
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 22
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                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/09/963,803 CURRENT FILING DATE: 2001-09-26 PRIOR APPLICATION NUMBER: FR 99/03925 PRIOR FILING DATE: 1999-03-29 PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello TITLE OF INVENTION: virus and cassava vein mosaic virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE:
                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                    FEATURE:
OTHER INFORMATION: promoter MPr1165
                                                                                                                                                                NAME/KEY: promoter LOCATION: (1)..(600)
258 GATTGATGTGATATCTCCACTGACGTAAGGGATGACGCATGCCAC 302
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                                                                 Conservative
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                                                                                 71.48; 100.08;
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Pred. No.
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Pred. No. 1.2e-
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                                                               . 1.3e-08; ches 0;
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LENGTH: 566
TYPE: DNA
ORGANISM: COIN
FEATURE:
NAME/KEY: DNA
LOCATION: (1)..(566)
OTHER INFORMATION: 5' genome+insert sequence
US-09-951-470-3
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                                                                                                                                               ; OTHER INFORMATION: Recombinant Cauliflower Mosaic Viral Promoter (CaMV35S) US-09-943-692-33
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US-09-943-692-33
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                                                                                          Query Match
Best Local Similarity
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SOFTWARE: PatentIn version 3.1

SEQ ID NO 33

LENGTH: 661
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Corn Event MON 810 and Compositions and Methods for Detection The FILE REFERENCE: 38-21 (5220)B CURRENT APPLICATION NUMBER: US/09/951,470 CURRENT FILING DATE: 2001-09-13 PRIOR APPLICATION NUMBER: 05/232,208 PRIOR APPLICATION NUMBER: 060/232,208 PRIOR APPLICATION NUMBER: 060/232,208 PRIOR FILING DATE: 2000-09-13 NUMBER: 05 SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                            CURRENT FILING DATE: 2001-08-31 PRIOR APPLICATION NUMBER: 09/027,998 PRIOR FILING DATE: 1998-02-23
                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/943,692 CURRENT FILING DATE: 2001-08-31
                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: FUCHS, ROY L.
APPLICANT: LAVRIK, PAUL B.
APPLICANT: MCPHERSON, SYLVIA A.
APPLICANT: PERLAK, FREDERICK J.
TITLE OF INVENTION: COLEOPTERAN TOXIN PROTEINS OF BACILLUS THURINGIENSIS
FILE REFERENCE: MOBT:195--1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: FISCHHOFF, DAVID A.
                                                                                                                                                                                   ORGANISM: Artificial Sequence FEATURE:
435 CAAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 479
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12 CTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 56
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                                                                           44; Conservative
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                                                                                          68.9%;
97.8%;
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97.88;

    Mismatches

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                                                                                        Score 43.4; DB 10 Pred. No. 6.1e-08;
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Pred. No. 5.9e-08;
                                                                         Mismatches
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                   Arabidopsis Genome 
Unpublished (2001)
                                                                                                                        Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnollophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis. 1 (bases 1 to 193)
1 (bases 1 to 193)
Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,P., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R., Gadrinab,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L., Shinn,I., Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                  Class: TDNA tagged.
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Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Salk Institute Genomic Analysis Laboratory (SIGnAL) The Salk Institute for Biological Studies
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/db_xref="taxon:3702"
/clone="SALK_045097.54.25.x"
/clone_11b="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
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                                                                                                                                                      153 bp DNA linear GS SALK_040762 Arabidopsis thaliana TDNA insertion lines thaliana genomic clone SALK_040762, DNA sequence.
BH619283
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Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Predni, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92
Tel: 858 453 4100 x1752
Fax: 858 558 6379
вн619283.1
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Unpublished (2001)
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44; Conservative
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/strain="Columbia /
/strain="SALK_047736.48.15.x"
/clone="SALK_047736.48.15.x"
/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/clone_lib="Arabidopsis thaliana TDNA insertion lines |
/note="PCR was performed on Arabidopsis thaliana lines |
each of which contains one or more TDNA insertion |
elements. The resultant fragment for each lines was |
directly sequenced to determine the genomic sequence at |
the site of insertion. Details of the protocols used can |
be found at http://signal.salk.edu/tdna_protocols.html"
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                 Arabidopsis Genome
Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
                                                                                                                                                                                                                           Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Predni, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in
                                                                                                                       ,C., Jeske,A., Karnes,M., Kim,C.J., Parl
, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
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Tel: 858 453 4100 x1752
Fax: 858 558 6379
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44; Conservative
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/note="PCR was performed on Arabidopsis thallana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
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/db_xref="taxon:3702"
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/<train="Columbia 0"
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Parker,H., Prednis,L.,
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Fax: 858 558 6379
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Tel: 858 453 4100 x1752
Fax: 858 558 6379
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Contact: Joseph R. Ecker
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L
, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., C., Jeske,A., Karnes,M., Kim,C.J., Parker,H.,
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The Salk Institute for Biological Studies
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Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
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/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/clone_lib="Arabidopsis thaliana lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 37 c 33 g 36 t
                /organism="Arabidopsis thaliana"
/strain="Columbia 0"
/db_xref="ftaxon:3702"
/clone="SALK_045100.51.10.x"
/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines each of which contains one or more TDNA insertion elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
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/db_xref="taxon:3702"
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/strain="Columbia 0"
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Pred. No. 4.2e-05;
0; Mismatches 1
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                                                                                                                          Local
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Salk Institute Genomic Analysis Laboratory (SIGNAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
1 (bases 1 to 165)
Alongo. Ture Train - The Tra
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165 bp DNA linear GSS 27-FEB-2002 SALK_016522.55.50.x Arabidopsis thaliana TDNA insertion lines Arabidopsis thaliana genomic clone SALK_016522.55.50.x, DNA
     BH746616
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                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                       /clone_lib="Arabidopsis thallana TDNA insertion lines"
/note="FCR was performed on Arabidopsis thallana lines
each of which contains one or more TDNA insertion
elements The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
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/db_xref="taxon:3702"
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/strain="Columbia 0"
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                                                                                                                                                                                                                               Score 43.4; DB 17;
Pred. No. 4.3e-05;
0; Mismatches 1;
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Contact: Joseph R. Ecker
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                                                                                       (bases 1 to 190)
onso,J.M., Leisse,T.J.,
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97.8%;
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12 CTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 56
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                   ALONSO,J.M., Leisse,T.J., Barajas,P., Chen,H.,,C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Zimmerman,J. and Ecker,J.R.
                                                                                                                                                Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheop
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
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/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 43 c 40 g 38 t
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/clone="SALK_045268.54.50.x"
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Library of Insertion
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Pred. No. 4.4e-05;
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                                                                                            Cheuk, R.,
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Prednis, L., Shinn, I
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Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                Unpublished (2001)
Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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                                                                                                                   Class: TDNA tagged
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  /organism-"Arabidopsis thaliana"
/strain="Columbia 0"
/db_xref="taxon:3702"
/clone="SALK_035354.55.75.x"
                                                                                                 Location/Qualifiers
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/clone_lib="Arabidopsis thaliana TDNA insertion lines"
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97.8%;
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Tel: 858 453 4100 x1752
Fax: 858 558 6379
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Salk Institute Genomic Analysis Laboratory (SIGnAL) The Salk Institute for Biological Studies 10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
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Alonso,J.M., Leisse,T.J., Barajas,P., Chen,H., Cheuk,R.,
C., Jeske,A., Karnes,M., Kim,C.J., Parker,H., Prednis,L
Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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Unpublished (2001)
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                                                                            /clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="FCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
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                     Score 43.4; DB 17
Pred. No. 4.8e-05;
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Pred. No. 4.8e-05;
Mismatches
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Prednis, L., Shinn,
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RESULT 13
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Best Local :
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Arabidopsis thaliana
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots;
Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
1. (bases 1 to 220)
                                                                                                                                                                                                               Arabidopsis
                                                                                                                                                                                                                                      BH747438
SALK_017469.
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The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
Email: ecker@salk.edu
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Unpublished (2001)
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A Sequence-Indexed Library of Insertion Mutations in the
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/clone_lib="Arabidopsis thaliana TDNA insertion lines"
/note="PCR was performed on Arabidopsis thaliana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"

57 c 50 g 46 t
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1. .219
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/strain="Columbia 0"
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97.88;
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Pred. No. 4.8e-05;
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Best Local Similarity
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                                                                                                                                                                855 California Ave, Palo Al
Tel: 650 723 2227
Fax: 650 725 8221
Email: walbot@stanford.edu
                                                                                                      Reverse complemented post-ligation Plate: 1008026 row: 10
                                                                                                                                                                                                                                                        Stanford University
                                                                                                                                                                                                                                                                                                 Unpublished (2001)
Contact: Walbot V
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Spermatophyta; Magnoliophyta; Liliopsida;
clade; Panicoldeae; Andropogoneae; Zea.
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, Zimmerman,J. and Ecker,J.R.
A Sequence-Indexed Library of Insertion Mutations in the
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                                                                                                                                               Very probable ligation site of
                                                                                                                                                                                                                                                                         Department of Biological
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/note="PCR was performed on Arabidopsis thallana lines
each of which contains one or more TDNA insertion
elements. The resultant fragment for each line was
directly sequenced to determine the genomic sequence at
the site of insertion. Details of the protocols used can
be found at http://signal.salk.edu/tdna_protocols.html"
a 55 c 48 g 50 t
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/strain="Columbia 0"
/organism="Zea mays"
/cultivar="mixed background W23/A188/B73'
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/clone="SALK_017469.51.60.x"
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Pred. No. 4.8e-05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Joseph R. Ecker
Salk Institute Genomic Analysis Laboratory (SIGnAL)
The Salk Institute for Biological Studies
10010 N. Torrey Pines Road, La Jolla, CA 92037, USA
Tel: 858 453 4100 x1752
Fax: 858 558 6379
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ecker@salk.edu
/strain="Columbia 0"
/db_xref="taxon:3702"
/db_xref="taxon:3702"
/clone="sakk_042817.55.00.x"
/clone=lib="Arabidopsis thaliana TDNA insertion lines"
/clone_lib="Arabidopsis thaliana TDNA insertion lines each of which contains one or more TDNA insertion elements. The resultant fragment for each line was directly sequenced to determine the genomic sequence at the site of insertion. Details of the protocols used can be found at http://signal.salk.edu/tdna_protocols.html"
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/strain="Columbia 0"
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/clone_lib="1008 - Re
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Pred. No. 4.8e-05;
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ACCESSION VERSION KEYWORDS

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RESULT 15 BH746474 LOCUS

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Matches

Search completed: May 24, 2003, 15:33:26	Qy 12 CTAGT	Query Match 68.9%; Score 43.4; DB 17; Length Best Local Similarity 97.8%; Pred. No. 4.9e-05; Matches 44; Conservative 0; Mismatches 1; Indels	BASE COUNT ORIGIN
d: May 24,	12 CTAGTGGATTGATGTGATATCTCCACTGACGTAAGGGATGACGCA 56	nilarity Conserva	68 a
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score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

RESULT 1 AX036743 LOCUS DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM REFERENCE AUTHORS TITLE Result 0000 000 O ŏ Score AX036743 Sequence 9 AX036743 AX036743.1 artificial sequences. 1 (bases 1 to 60) Rance,I., Theisen,M. and Gruber,V. Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 9 05-OCT-2000; synthetic construct. synthetic construct Match 11000 1036 8159 8340 8340 12241 143649 167852 176454 177876 1778 Length 9285 15077 60 bp from Patent WO0058485. GI:11226252 밁 AX088390 AX014764 AX088391 AX088393 AX088393 AX088393 AX088393 AX088393 AX014765 CVU20341 CVU209751 AX329231 AX339231 AX339231 AX339231 AX412168 ACC120006 ACC12006 ACC111873 ACC124942 ACC011387 ACC124942 ACC11873 ACC1 AX036759 AX036758 AX036758 AX036757 AX036736 AX036736 AX088388 AX202413 AX036755 AX036753 AX036754 AX036756 AX036743 AX036737 AX093047 AX093052 AX036741 ALIGNMENTS DNA linear AX014765 Sequence U20341 Cassava vei U59751 Cassava vei AX329231 Sequence AX412168 Sequence AX412168 Sequence AC120006 Mus muscu AC072061 Homo sapi AC111873 Rattus no AC124942 Rattus no AC124943 Rattus no AC095908 Rattus no AC011323 Homo sapi AL672033 Mus muscu AC098477 Homo sapi AL646019 Human DNA AC115910 Mus muscu AX036740 AX036759 AX036758 AX036757 AX036757 AX036736 AX088388 AX202413 AX088391 AX088391 AX088392 AX088393 AX088393 AX088393 AX014765 U20341 C AX088390 AX014764 Description PAT 16-NOV-2000 Sequence Human DNA Human DNA Sequence

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l (bases 1 to 348)

1 (bases 1 to 348)

2 (chimeric expression promoters originating from (mottle virus and cassava vein mosaic virus patent: WO 0058485-A 4 05-OCT-2000;

MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);

GRUBER VERONIQUE (FR)

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                           1 (bases 1 to 393)
1 (bases 1 to 393)
Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating fr
mottle virus and cassava vein mosaic virus
Patent: WO 0058465-A 19 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
LOCATION/Qualifiers
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Rance, I., Thelsen, M. and Gruber, V.
Chimeric expression promoters originating mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 21 05-OCT-2000;
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AX036756
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AX036754
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Best Local Similarity
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Best Local
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COUNT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 00548485-A 20 05-OCT-2000;
MERISTEM THERAFEUTICS (FR); RANCE IANN (FR); GRUBER VERONIQUE (FR);
                                                                                            Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 22 05-OCT-2000;
MERISTEM THERAPBUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                       600 bp
Sequence 22 from Patent W00058485.
AX036756
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1 (Dases 1 to 462)
Rance, I., Theisen, M. and Gruber, V.
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                                                                                                                                                                              artificial sequences.
                                                                                                                                                                                            synthetic construct
                                                                                                                                                                                                       synthetic construct.
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                                /organism="synthetic construct"
/db_xref="taxon:32630"
/note="promoter MPr1165"
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/db_xref="taxon:32630"
/note="promoter MPr1163"
                                                                                 Location/Qualifiers
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87 c
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Pred. No. 3.4e-10;
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WO0058485.
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REFERENCE
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VERSION
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AX093052
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ORGANISM
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KEYWORDS
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AX093047
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Best Local Similarity
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Best Local
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AX093047
                                                                                                                                                                                                                                                                                                    15077 bp
Sequence 57 from Patent WO0118192.
AX093052
                                                                                                                                                                                  artificial sequences.

1 (bases 1 to 15077)

Gruber, V. and Comeau, D.

Synthetic vectors, transgenic plants
                                                                                                                                                                                                                                               synthetic construct. synthetic construct
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Gruber, V. and Comeau, D.
Synthetic vectors, transgenic plants containing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                synthetic construct 
synthetic construct
                                                                                                                              MERISTEM THERAPEUTICS (FR)
Location/Qualifiers
                                                                                                                                                         for obtaining them Patent: WO 0118192-A 57 15-MAR-2001;
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ilarity 100.0%;
Conservative 0
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/note="pMRT1342 results from the replacement of the expression cassette ep35S-gus-polyA35S from pMRT1335 by the expression cassette L5-gus-polyA35S isolated from pMRT1336"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /note="pMRT1336 results from the insertion into pMRT1196 of the promoter MPT1165 isolated from plasmid pMRT1322 as described in PCT patent application PCT/IB00/00370" a 2252 c 2506 g 2087 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     52
                                                                    /organism="synthetic construct"
/db_xref="taxon:32630"
/note="pMRT1342"
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/db_xref="taxon:32630"
/note="pMRT1336"
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Pred. No. 3.2e-10;
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Pred. No. 3.4e-10;
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W00118192
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BASE COUNT
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AX036739
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AX036741
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Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   synthetic construct
artificial sequences.

1 (bases 1 to 301)
Rance,I., Theisen,M. and Gruber,V.
Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus
Patent: WO 0059485-A 7 05-OCT-2000;
MERISTEM THERAPEUTICS (FR); RANCE IANN (FR);
GRUBER VERONIQUE (FR);
                                                                                                  Rance, I., Theisen, M. and Gruber, V. Chimeric expression promoters originating from mottle virus and cassava vein mosaic virus Patent: WO 0058485-A 5 05-0CT-2000; MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                               AX036739 371 bp
Sequence 5 from Patent WO0058485.
AX036739 AX036739.1 GI:11226248
                                                                                                                                                                                                                    synthetic construct.
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AX036741
                                                                                                                                                                                            artificial sequences.
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                                                                                                                                                                                                           synthetic construct
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56; Conservative
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                                    /organism="synthetic con./db_xref="taxon:32630"
/note="promoter MPr1146"
                                                                                      Location/Qualifiers
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/db_xref="taxon:32630"
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Pred. No. 1.3e-08;
0; Mismatches 1;
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Pred. No. 3.2e-10;
"Hamatches 0;
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SOURCE
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DEFINITION
ACCESSION
VERSION
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AX036759
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AUTHORS
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Query Match
Best Local Similarity 98.7
Matches 56; Conservative
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Sequence 25
AX036759
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Sequence 6
AX036740
                                                                                                                                                   artificial sequences.

1 (bases 1 to 472)
Rance, I., Theisen, M. and Gruber, V.
Chimeric expression promoters originating i
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 25 05-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 (bases 1 to 398)
Rance, I., TheLsen, M. and Gruber, V.
Chimeric expression promoters originating
mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 6 05-OCT-2000;
                                                                                                                                                                                                                                                                                                                                                                                     Similarity
56; Conserv
                                                                                                                     MERISTEM THERAPEUTICS (FR); RANCE; GRUBER VERONIQUE (FR)
Location/Qualifiers
                                                                                                                                                                                                                        synthetic
                                                                                                                                                                                                                                synthetic construct.
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/db_xref="taxon:32630"
/note="promoter MPr1147"
                                                                                      organism="synthetic construct"/db_xref="taxon:32630"
                                                                            note-"promoter MPr1169"
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from Patent WO0058485
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AUTHORS
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AX036757
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AX036758
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                                            Query Match
Best Local Similarity
Matches 56; Conserv
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Best Local Similarity
Matches 56; Conserva
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1 (bases 1 to 541)

Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus
Patent: WO 0058485-A 24 05-CCT-2000;

MERISTEM THERAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR)
                                                                                                                                                                                                            artificial sequences.

1 (bases 1 to 604)

Rance, I., Theisen, M. and Gruber, V.

Chimeric expression promoters originating from commelina yellow mottle virus and cassava vein mosaic virus

Patent: WO 0058485-A 23 05-OCT-2000;

MERISTEM THERRAPEUTICS (FR); RANCE IANN (FR); THEISEN MANFRED (FR);

GRUBER VERONIQUE (FR)
                                                                                                                                                                                                                                                                                                                                                                              604 bp
Sequence 23 from Patent WO0058485.
AX036757.
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Sequence 24 from Patent WO0058485.
AX036758
                                                                                                                                                                                                                                                                                                                           synthetic construct
                                                                                                                                                                                                                                                                                                                                       synthetic construct.
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nilarity 98.2%;
Conservative
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/db_xref="taxon:32630"
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/db_xref="taxon:32630"
/note="promoter MPr1168"
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                                                                                                                                         note-"promoter MPr1167"
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116 c
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98.2%;
                                          Score 55.4; DB 6; Pred. No. 1.3e-08; 0; Mismatches 1;
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Pred. No. 1.3e-08;
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Db 160 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 216

Search completed: May 24, 2003, 14:52:19
Job time: 340.194 secs

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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Result
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                                                                                                                                                                                                                                        Score
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /SIDS2/gcgdata/geneseq/geneseqn-embl/NA1980.
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1981.
/SIDS2/gcgdata/geneseg/geneseqn-embl/NA1983.
/SIDS2/gcgdata/geneseq/geneseqn-embl/NA1983.
                                                                                                                                                                                                                                        Length
         317
318
348
392
393
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301
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         Ä
       AAA96837
AAA96838
AAA96858
AAA96858
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AAA96841
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1304.896 Million cell updates/sec
Nucleotide sequenc
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Arabidopsis thalia	AAC45506 AAC40898	21	2525 2547	42.0	25. 25. 2	4 τυ
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70		19	305		7.	_
	AAD	24	12241	.7		0
		24	12241	.7	40.4	9
pAGI4002 binary ve		24	8340	.7	40.4	8
Binary vector		24	8340	.7	40.4	7
4-Hydroxyphenylpyr		24	8187	.7	40.4	6
4-Hydroxyphenylpy		24	4677	.7	40.4	G
4-Hydroxyphenylpyr		24	1839	7.	40.4	4
Nucleotide sequenc		22	931	7.	40.4	ω
Nucleotide sequenc		22	931	7	40.4	N
Nucleotide sequenc		22	931	7.	40.4	_
		22	857	7.	40.4	0
Nucleotide sequenc		22	853	7.	40.4	9
Nucleotide sequenc		22	593	7.	40.4	æ
<		22	532	7.	40.4	7
CsVMV promoter		19	526	7.	40.4	6
CsVMV promoter		19	524	7.	40.4	5
Nucleotide sequenc		22	515	.7	40.4	4
Promoter from		21	515	.7	40.4	ω
CsVMV promoter		19	491	.7	40.4	N
		19	476	7.	40.4	_
•		19	468	7.	40.4	0
CsVMV promoter		19	458	.7	40.4	9
		19	441	.7	40.4	æ
		19	418	.7	40.4	7
CsVMV promoter		19	411	7.	40.4	σ
CsVMV promoter		19	408	.7	40.4	51
CsVMV promoter		19	392	7.	40.4	4
		21	Ò		55.4	ω
	AAA9	21	541	٢		N
	AAA968	21	7	92.3	55.4	_
Nucleotide sequenc	AAA96840	21	398	2	55.4	0

a

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ALIGNMENTS

RESULT 1
AAA96843
ID AAAS promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression pro plant green tissue expression promoter; Cassava vein mosaic 29-MAR-1999; Directional desoxynucleotide building block \$2. AAA96843 standard; DNA; 60 BP Rance I, (MERI-) MERISTEM THERAPEUTICS. 29-MAR-2000; 2000WO-IB00370 05-OCT-2000 WO200058485-A1 Synthetic transgenic plant; ss. 19-FEB-2001 (first entry) AAA96843; Gruber V, 99FR-0003925 Theisen M;

ion promoter;
mosaic virus;

Chimeric expression promoter for transgenic plant production, comprises

WPI; 2000-647238/62

0

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RESULT 2
AAA96837
ID AAA9
XX AAA9
XX AAA9
XX Prom
XW Chir
XX Prom
XW Chir
XX Prom
XW Chir
XX Prom
XW Chir
XX Prom
XX Prom
PN WO2:
YX 29:
YX 29:
YX 29:
YX AMA9
PR 29:
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Best
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with sequ
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Chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       chimeric
plant gre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        block, which was used to construct chimeric promoters of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transcenic regions.
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The present sequence represents a chimeric promoter of the in
The specification describes chimeric expression promoters. Th
chimeric promoters comprise a nucleic acid sequence which is
from a first plant promoter, in which a plant vascular expres
                                                                              Claim
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                                                                                                                                                                                                                                                                                    29-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                 05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      19-FEB-2001
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                                                                                                                                                                                                                                                     (MERI-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          transgenic
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                                                                                                                           ence from
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                                                                                                                                                                                         2000-647238/62
                                                                              5:
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                                                                                                                                                                                                                                                     MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
                                                                            Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8
                                                                                                                                                          expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression promoter;
een tissue expression
                                                                                                                                                                                                                     Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                           Commelina yellow mottle virus. Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sequence of chimeric expression promoter MPr1116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP;
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                                                                                                                                                                                                                                                                                                                2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                            81;
                                                                                                                        ession promoter for transgenic plant production, compris
promoter comprising vascular expression region replaced
from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 promoter comprising vascular expression region replaced from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                   99FR-0003925
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                                                                          91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         chimera;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DNA;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      plant vascular expression promoter; Cassava vein mos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    G; 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mismatches
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J. 3.2e-12;
O;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       mosaic virus
 expression
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                              invention These
                derived
                                                                                                                                                     comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    building
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Query Match

Sequence

348

BP;

A;

78

ç; 60; 84

H

0 21;

100.0%; 70 ·C;

Score

BB

Length

intergenic regions. transgenic plants.

The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful

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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local
                                                                                                                                                                                Chimeric expression promoter for transgenic plant production, comprises sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.
                                                                                                                                                    Claim
                                                                                                                                                                                                                                                                                                                                                                                Chimeric - Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                               chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.
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                                                                                                                                                                                                                                                  Rance
                                                                                                                                                                                                                                                                                                                                      05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide sequence of chimeric expression promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAA96838
                                                                                                                                                                                                                                                                      (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                             29-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                                           WO200058485-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                               Promoter; intergenic region; Commelina yellow mottle
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA
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                                                                                                                                                    5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                60; Conserv
                                                                                                                                                   Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          317
                                                                                                                                                                                                                                                  Gruber
                                                                                                                                                                                                                                                                                                                                                                               Cassava
                                                                                                                                                                                                                                                                                                                                                                                          Commelina
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llarity 100.0%;
Conservative C
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                                                                                                                                                91pp;
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                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                               mosaic
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                virus;
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promoter MPr1162

mosaic virus;

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RESULT 5
AAA96853
ID AAA9
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AAA96855
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                                                                                                                    Matches
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                                                                                                                                                                                                      The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                                                                                                        Chimeric
sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI;
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 AAA96853 standard;
                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                       Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            chimeric expression promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Nucleotide sequence of chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19-FEB-2001
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                                                                  75
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                                                                                                                                             Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       _
                                                                                                                                                                                              producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 60
                                                                                                                                                                                                                                                                                                                                                                                             sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                               2000-647238/62.
                                                               AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ter; intergenic region; Commelina yellow mottle virus; ric expression promoter; plant vascular expression promoter; green tissue expression promoter; Cassava vein mosaic virus;
                                                                                                                                                                                                                                                                                                                                                       5.
                                                                                                                               Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MERISTEM THERAPEUTICS
                                                                                                                                                                        392
                                                                                                                                                                                                                                                                                                                                                                                        expression promoter for transgenic plant production, compristion promoter comprising vascular expression region replaced uence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                      Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Commelina yellow mottle virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   plant;
                                                                                                                  100.0%; llarity 100.0%; Conservative 0;
                                                                                                                                                                     BP;
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                                                                                                                                                                                                                                                                                                                                                       66;
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                                                                                                                                                                     127 A; 80 C;
                                                                                                                                                                                                                                                                                                                                                      91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     chimera;
  DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Theisen M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
                                                                                                                               Score 60;
Pred. No.
                                                                                                                                                                     87 G; 98 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 4.7e-12; Mismatches 0;
                                                                                                                   Mismatches
                                                                                                                               DB 21;
4.8e-12;
                                                                                                                                                                     0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  promoter MPr1164
                                                                                                                  0;
                                                                                                                                          Length
                                                                                                                   Indels
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                                                                                                                                           392;
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                                                                                                                                                                                                                                                                                                                                                                                                                    comprises
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                                                                                                                Gaps
                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0
Q
                                                                                                                                                                                                                                                                                                              RESULT
                                                                                                                                                                                                          В
                                                                                                                                                                                                                                                                       Query Match
Best Local
                                                                                                                                                                                                                                                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chimeric expression promoter for transgenic plant production, compris-
sequence from promoter comprising vascular expression region replaced
with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chimeric - Commelir
Chimeric - Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           chimeric expression promoter; plant vascular explant green tissue expression promoter; Cassava transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                              Sequence 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 2000-647238/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            29-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       19-FEB-2001
                                                                 Nucleotide
                                                                                                                                           AAA96854 standard; DNA; 462 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MERI-) MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29-MAR-1999;
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                                                                                                                                                                                                          75
                                                                                                                                                                                                                                    μ
                                                                                                                                                                                                                                                                                                                                       producing
                                                                                                                                                                                                          AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 60
                                                                                                                                                                                                                                                                        Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          intergenic region; Commelina yellow mottle virus; expression promoter; plant vascular expression pro en tissue expression promoter; Cassava vein mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Commelina yellow mottle virus
                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                              BP; 128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2000WO-IB00370
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                                                                                        (first entry)
                                                                                                                                                                                                                                                                                                                                       transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99FR-0003925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  vein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              of chimeric expression
                                                                                                                                                                                                                                                                       100.0%;
                                                                                                                                                                                                                                                                                                             A; 75 C; 93 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Theisen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         mosaic
                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 X
                                                                                                                                                                                                                                                           Score 60; DB Pred. No. 4.8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus.
                                                                                                                                                                                                                                                                                                              97 T; 0 other;
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Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus;
                                                                                                                                                                                                 sequence of chimeric expression
                                                                                                                                                                                                 promoter MPr1163
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4.8e-12; BB

0;

Gaps

0

21;

Length Indels

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RESULT 7
AAA96856
ID AAA9
AC AAA9
AC AAA9
AC AAA9
XX Nucl
XX Prom
KW Prom
KW Chim
KW Plan
KW Plan
KW Chim
XX C
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Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mossic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                               Chimeric - Chimeric -
                                                                                                                                                                                                                                                                                                                                         Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chimeric expression promoter for transgenic plant production, compris sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Chimeric -
     29-MAR-1999;
                                                          29-MAR-2000;
                                                                                                                  05-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 5; Page 86; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              29-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 462
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
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                                                                                                                                                                                                                            Commelina yellow mottle virus Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                     plant;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence of chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   BP; 148 A; 87 C; 111 G; 116 T; 0 other;
                                                          2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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     99FR-0003925
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                                                                                                                                                                                                                                                                                                                     chimera;
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                                                                                                                                                                                                                                                                                                                        88
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            promoter
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RESULT 8
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chimeric expression promoter for transgenic plant production, comprise sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                     05-OCT-2000
                                                                                                                                                                                                                                   Chimeric -
                                                                                                                                                                                                                                                                                    chimeric expression promoter; plant vascular expression plant green tissue expression promoter; Cassava vein mos transgenic plant; chimera; ss.
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                                                                                                   29-MAR-1999;
                                                                                                                                  29-MAR-2000; 2000WO-IB00370
                                                                                                                                                                                                                                                                                                                                     Promoter; intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                  Nucleotide sequence of chimeric expression promoter MPr1154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAA96841 standard;
                                                                (MERI-) MERISTEM THERAPEUTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      l Similarity
60; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page
                               Gruber
                                                                                                                                                                                                                                   Commelina yellow mottle virus. Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence represents a chimeric promoter of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP; 188 A; 111 C;
                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          86-87;
                                                                                                 99FR-0003925
                             <
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA;
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                               Theisen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91pp;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    147 G; 154 T;
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                                                                                                                                                                                                                                                                                                                   promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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0

Chimeric expression promoter for transgenic plant production, sequence from promoter comprising vascular expression region; with sequence from promoter comprising green tissue expression

expression

n replaced on region

0;

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RESULT 9
AAA96839
ID AAA9
XX AAA9
XX AAA9
XX Pron
DT 19-F
XX Pron
KW Chin
KW Plan
KW Chin
KW HTGH
XX Pron
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KW HTGH
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Chi
PR 29-I
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Best Local
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                  The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from
                                                                                                                                                                                                                                                                                                                                                                    Chimeric
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Chimeric -
Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus
                                                                                                                                                                                                                                      Claim 5;
                                                                                                                                                                                                                                                                                                                                         sequence from
                                                                                                                                                                                                                                                                                                                                                                                                                       WPI;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-MAR-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200058485-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nucleotide sequence of chimeric expression promoter MPr1146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19-FEB-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              transgenic plant; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         sequence
                                                                                                                                                                                                                                                                                                                                                                                                                    2000-647238/62
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    301
                                                                                                                                                                                                                                                                                                            expression promoter for transgenic plant production, compris
from promoter comprising vascular expression region replaced
uence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Page
                                                                                                                                                                                                                                      Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gruber
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Commelina yellow mottle virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2000WO-IB00370
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                                                                                                                                                                                                                                      81;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99FR-0003925
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ۲,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98 A; 54 C;
                                                                                                                                                                                                                                 91pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              91pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DNA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92.3%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   371
                                                                                                                                                                                                                                   English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ВP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 55.4;
Pred. No. 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       No. 2e-10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ВG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       vein mosaic virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               301;
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originates
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CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA

Query Match Best Local

Similarity

92.3%; 98.2%;

Conservative

0

Score 55.4; D Pred. No. 2.2e 0; Mismatches

,2e-10

Indels Length

0;

Gaps

0

60

В 0 other;

21;

Sequence

398

B₽;

128 A; 80 C;

93

G;

97 T;

Matches

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RESULT 10
AAA96840
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Best Local
                                   chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originate from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                     Chimeric sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                 29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chimeric - Chimeric -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAA96840 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                    The present sequence represents a chimeric promoter of the invention The specification describes chimeric expression promoters. These
                                                                                                                                                                                              Claim
                                                                                                                                                                                                                                                                                                                           kance
                                                                                                                                                                                                                                                                                                                                                                                                             29-MAR-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAA96840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                     (MERI-) MERISTEM THERAPEUTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  130 CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4
                          producing transgenic plants.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          producing
                                                                                                                                                                                                                                    eric expression promoter for transgenic plant production, comprisence from promoter comprising vascular expression region replaced sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              371
                                                                                                                                                                                              Page
                                                                                                                                                                                                                                                                                                                        Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Commelina yellow mottle virus. Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       plant;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           sequence of chimeric expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              вр;
                                                                                                                                                                                                                                                                                                                                                                                                             2000WO-IB00370
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transgenic plants.
                                                                                                                                                                                              82;
                                                                                                                                                                                                                                                                                                                                                                                 99FR-0003925.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              122 A; 68 C; 89
                                                                                                                                                                                              91pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       e expression chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     entry)
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98.2%;
                                                                                                                                                                                                                                                                                                                          Theisen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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Pred. No. 2.1e
0; Mismatches
                                                                                                                                                                                                                                                                                                                           X
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              92 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .1e-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           promoter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21;
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                                                    promoter originates promoters are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
                                                                                                                                                                                                                                                                  comprises
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RESULT 12
AAA96858
ID AAA96
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AC AAA96
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AAA96859
ID AAA96
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                                                                                                                        Query Match
Best Local S
Matches 56
                                                                                                                                                                                       The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from commelina yellow mottle virus, and the second plant promoters are derived from intergenic regions. The chimeric promoters are useful
                                                                                                                                                                                                                                                                                                                       Chimeric expression promoter for transgenic plant production, compris sequence from promoter comprising vascular expression region replaced with sequence from promoter comprising green tissue expression region
                                                                                                                                                                                                                                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         chimeric expression promoter; plant green tissue expression transgenic plant; chimera; ss.
                   AAA96858;
                                     AAA96858 standard;
                                                                                                                                                                                                                                                                                              Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chimeric -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Nucleotide
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19-FEB-2001
                                                                                                                                                             Sequence 472
                                                                                                                                                                                                                                                                                                                                                                                 Rance
                                                                                                                                                                                                                                                                                                                                                                                                  (MERI-)
                                                                                                                                                                                                                                                                                                                                                                                                                     29-MAR-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                       29-MAR-2000;
                                                                                                                                                                                for producing transgenic plants.
                                                                                    160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            160
                                                                                                      4 CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 60
                                                                                    CATGCCACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA
                                                                                                                                                                                                                                                                                                                                                              2000-647238/62.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity 98.
                                                                                                                                                                                                                                                                                                                                                                                                  MERISTEM THERAPEUTICS
                                                                                                                                                                                                                                                                                              Page 88;
                                                                                                                                                                                                                                                                                                                                                                                Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     intergenic region; Commelina yellow mottle virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Commelina yellow mottle virus. Cassava vein mosaic virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       sequence
                                                                                                                                                             BP; 149 A; 92 C;
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(first
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                                                                                                                                                                                                                                                                                              91pp; English
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entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of chimeric expression promoter MPr1169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        entry)
                                                                                                                                92.3%;
98.2%;
                                                                                                                                                                                                                                                                                                                                                                                Theisen
                                     541
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                                     ВP
                                                                                                                        Score 55.4; D
Pred. No. 2.2e
0; Mismatches
                                                                                                                         0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 plant vascular expression promoter; promoter; Cassava vein mosaic virus
                                                                                                                                                              112 G;
                                                                                                                                                             119 T; 0 other;
                                                                                                                                2e-10;
                                                                                                                                          DB 21;
                                                                                                                                          Length
                                                                                                                         Indels
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Best Local S
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Chimeric -
                 Chimeric
                                                                                    Nucleotide
                                                                                                      19-FEB-2001
                                                                                                                         AAA96857
                                                                                                                                          AAA96857 standard;
                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                 Claim
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Nucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MERI-) MERISTEM THERAPEUTICS.
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                                                                                                                                                                                                                                       Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          intergenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cassava
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             plant;
                  Commelina yellow mottle virus
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chimeric expression promoter; plant vasc plant green tissue expression promoter;
                                                                                  of chimeric expression
                                             region;
                                             Commelina
                        Commelina yellow plant vascular
      cular expression promoter;
Cassava vein mosaic virus;
                                                                                promoter
                                             mottle
                                                                                    MPr1168
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Commelina yellow mottle virus chimera; vein mosaic 88 virus

2000WO-IB00370 99FR-0003925.

Theisen M;

Chimeric expression promoter for transgenic plant production, esequence from promoter comprising vascular expression region rewith sequence from promoter comprising green tissue expression

5; Page 87-88; 91pp; English.

The present sequence represents a chimeric promoter of the invention. The specification describes chimeric expression promoters. These chimeric promoters comprise a nucleic acid sequence which is derived from a first plant promoter, in which a plant vascular expression promoter region is replaced with a nucleic acid sequence derived from a second plant promoter comprising a plant green tissue expression promoter region. Preferably, the first plant promoter originates from Commelina yellow mottle virus, and the second plant promoter originates from the Cassava vein mosaic virus. Especially, the promoters are derived from intergenic regions. The chimeric promoters are useful for producing transgenic plants.

541 BP; 169 A; 104 C; 130 G; 138 7; 0 other;

Conservative 92.3%; 0 Score 55.4; Pred. No. 2 Mismatches DB 21; .3e-10; Length 541; 0 Gaps 0

DNA; 604 ΒP

(first entry)

sequence of chimeric expression promoter

Promoter; intergenic region; Commelina yellow mottle virus; chimeric expression promoter; plant vascular expression promoter; plant green tissue expression promoter; Cassava vein mosaic virus; transgenic plant; chimera; ss.

Cassava vein mosaic virus

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RESULT 14
AAV14019
ID AAV1
AC AAV1
XX AAV1
XX AAV1
XX Case
KW Case
XX Case
XX W097
PN W097
XX W097
XX 20-(
XX X X 20-(
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XX X X X X X X X X X X X X X X X X X 
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Best Local S
Matches 56
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                                                                            Beachy RN, De Kochko
                                                                                                                                                                                 20-JUN-1996;
                                                                                                                                                                                                                                 20-JUN-1997;
                                                                                                                                                                                                                                                                                   24-DEC-1997
                                                                                                                                                                                                                                                                                                                                     WO9748819-A1
                                                                                                                                                                                                                                                                                                                                                                                     Cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                     Cassava vein mosaic virus; CsVMV; promoter; cultivated crop; tissue-specific expression control; transgenic plant; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CSVMV promoter CVP1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAV14019 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                160
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     present sequence represents a chimeric promoter of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Page 87; 91pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gruber V,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
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                                                                                                                                                                            96US-0020129
                                                                                                                                                                                                                                 97WO-US10376.
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RESULT 15
AAV14031
ID AAV14
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Best Local
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         This sequence represents a cassava vein mosaic virus promoter, and is a nucleic acid molecule of the invention. The promoter is capable of initiating transcription of an operably linked heterologous nucleic acid sequence in a plant cell. The CsVMV promoters are active in both monocot and dicot plant species, and therefore can be readily applied to a variety of cultivated crops. Although generally constitutive, the derivative promoters include promoters that can regulate expression in a tissue-specific manner, and therefore are useful for controlling
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim
                                                                                                                                                     Claim
                                                                                                                                                                                                Cassava vein mosaic virus promoter - DNA sequences for producing transgen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         cassava vein mosaic virus; tissue-specific expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CSVMV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAV14031 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    phenotype(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cassava vein mosaic virus promoter - used to express heterologous DNA sequences for producing transgenic plants having altered
                                                                                                                                                                                                                                               WPI; 1998-063157/06
                                                                                                                                                                                                                                                                            Beachy RN,
                                                                                                                                                                                                                                                                                                                                         20-JUN-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cassava vein
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                                                                                                                                                                                                                                                                                                                                                                     20-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              143 GAATCTTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 184
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         promoter
                                                                                                                                                     2; Page 84-85; 115pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Page 74; 115pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
                                                                                                                                                                                                                                                                            De Kochko A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                               mosaic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 virus
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97.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CsVMV; promoter; cultivated crop;
control; transgenic plant; ss.
                                                                                                                                                     English.
                                                                                                                                                                                                transgenic
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Pred. No. 5
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                                                                                                                                                                                                used to express heterologous ic plants having altered
                                                                                                                                                                                                                                                                          Verdaguer
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heterologous genes

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CC phenotype.

Sequence 408 BP; 161 A; 61 C; 86 G; 100 T; 0 other;

ONEY MAICH
Best Local Similarity 97.6%; Pred. No. 5.1e-05;
MAICHES 41; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 19 CANTATCANGACGTANGCACTGACCACANCANGANAGAN 60

Db 222 CANTCTTHANGACCTGACCGACACACANCANGANAGAN 263

Search completed: May 24, 2003, 14:34:41

Job time: 104.548 secs
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Result
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB seq
                                                                                                        of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                     Issued_Patents_NA:*
1: /cgn2_6/ptodata/1,
2: /cgn2_6/ptodata/1,
3: /cgn2_6/ptodata/1,
4: /cgn2_6/ptodata/1,
5: /cgn2_6/ptodata/1,
6: /cgn2_6/ptodata/1,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    length: 0
length: 2000000000
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60
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        441362 seqs, 153338381 residues
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agccatgacactctgtgcga.....gacgacaacaatgaaaagaa
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                                                                                                                                                                                                                                                                                                                                                                                                   /cgn2_6/ptodata/1/ina/5A_COMB.seq:*
/cgn2_6/ptodata/1/ina/5B_COMB.seq:*
/cgn2_6/ptodata/1/ina/6A_COMB.seq:*
/cgn2_6/ptodata/1/ina/6B_COMB.seq:*
/cgn2_6/ptodata/1/ina/BCTUS_COMB.seq:*
/cgn2_6/ptodata/1/ina/backfiles1.seq:*
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13440
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                US-09-085-199B-3
US-09-472-971-7
US-09-474-971-7
US-09-404-879A-79
US-09-404-879A-79
US-09-404-879A-79
US-09-138-024-20
US-09-138-024-20
US-09-138-024-20
US-09-124-698-5
US-09-124-698-5
US-09-124-523-5
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740.804 Million cell updates/sec
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44, Appl
3, Appli
5, Appli
7, Appli
27, Appli
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391, Appli
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109, Appl
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375, Appli
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20, Appl
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ALI RESULT 1 US-08-961-527-168	0044444 0013848	28 29 31 31 31 33 33 34 38	
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œ ·	34.7 34.7 34.7 34.7 34.7 34.7	35.0 35.0 35.0 35.0 35.0 35.0 34.7 34.7	
	1524 1524 1524 2538 2912 3171 8878	3800 629 2102 2126 2258 2258 2258 2258 471 606 951	
	124111	40441054444	
ALIGNMENTS	US-07-752-420E-1 US-07-752-428C-1 US-07-752-428C-3 US-08-742-011-1 US-08-742-011-1 US-08-907-143-3 US-08-868-786-5 US-08-206-176-3	US-08-965-762-15 US-08-670-186-1 US-08-235-836C-65 US-08-235-836C-75 US-07-720-589-1 US-07-720-589-1 PCT-US92-05539-1 US-09-134-001C-395 US-09-134-001C-395 US-09-071-035-297 US-09-134-001C-1602	
	Sequence 1, Appli Sequence 1, Appli Sequence 3, Appli Sequence 1, Appli Sequence 3, Appli Sequence 5, Appli Sequence 3, Appli	Sequence 15, Appli Sequence 5, Appli Sequence 65, Appli Sequence 75, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 395, App Sequence 297, App Sequence 297, App	

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SEQUENCE CHARACTERISTICS:
LENGTH: 3025 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
US-08-961-527-168
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                                                               Best Loc
Matches
                                                                                              Query Match
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APPLICANT: Charles Kunsch
TITLE OF INVENTION: Strep
NUMBER OF SEQUENCES: 391
                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: BIOOKES, A. Anders
REGISTRATION NUMBER: 36,373
REFERENCE/DOCKET NUMBER: PB34
TELECOMMUNICATION INFORMATION:
TELEPHONE: (301) 309-8504
                                                                                                                                                                                                                                   TELEFAX: (301) 309-851 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: MSDO
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Diskette, 3.50 inch, 1.4Mb
2120 ATATTGAGGACACAATCACTGACTACCTTGAAAA 2156
                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
                 21 ATATTGAAGACGTAAGCACTGACGACAACAATGAAAA 57
                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Rockville
STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER:
                                                                              Local
                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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                                                               . Similarity 29; Conserv
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                                                               40.3%;
nilarity 78.4%;
Conservative
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                                                                                                                                                                                                                                                     309-8512
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                                                             Score 24.2; DB Pred. No. 4.9; O; Mismatches
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                                                                                              DB 4;
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                                                                                                Length 3025;
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                                                                                                                        Sequence 3, Application US/09085199B Patent No. 6235879
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Best Local S
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                                                                                                            GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (970) 668-2052 INFORMATION FOR SEQ ID NO:
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                                              APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                    FEATURE:
OTHER INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette,
   TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                      ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 32038
REFERENCE/DOCKET NUMBER: UBC.P-013US2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: WORDPERFECT CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                              APPLICANT:
                                                                                            APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: 96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                   546 AGTTTGTGCAAATGATGAAACTGAAAAGACTGAGAAGAAAACAGAAAAGAA 496
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                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: human
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TOPOLOGY: 11r
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Oppedahl & Larson
PO Box 5270
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Huq, A.H.M. Mahbubul
Chopra, Vikramjit Singh
                                          Hackam, Abigail
Huq, A.H.M. Mahbubul
Chopra, Vikramjit Singh
                                                                                                                                                                                                                                                                              Conservative
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                               Kalchman, Michael
                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
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                                                                                            Michael R.
                                                                                                                                                                                                                                                                                            39.7%;
Apoptosis Modulators
Huntington's Disease
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Pred..No. 7.3;
O; Mismatches
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 That Interact with the Gene
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; ORGANISM: Escherichia coli
US-09-472-971-5
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; OTHER INFORMATION:
US-09-085-199B-3
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Query Match
Best Local Similarity
Matches 32; Conserv
                                                                                                 SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 5
LENGTH: 1647
TYPE: DNA
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Best Local Similarity
                                                                                                                                                                                                                                                     SENERAL INFORMATION:
APPLICANT: SOGO, KAZUYO
APPLICANT: YANAGI, Hideki
APPLICANT: YURA, Takashi
TITLE OF INVENTION: TRIGGER FACTOR EXPRESSION PLASMIDS
FILE REFERENCE: 1422-409P
                                                                                                                                                                                    EARLIER APPLICATION NUMBER: JP10-372965 EARLIER FILING DATE: 1998-12-28
                                                                                                                                                                                                                       CURRENT FILING DATE:
                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/09/472,971
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (970) 668-2052
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                       NUMBER OF SEQ ID NOS:
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SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
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LENGTH: 4796
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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39.3%;
nilarity 69.6%;
Conservative
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Score 23.6; DI Pred. No. 7.1; 0; Mismatches
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                                 DB 4; Length 1647;
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RESULT 5 US-09-472-971-7 B 5

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1046 TCCGTCAGCAGATTGAAGAAGCAACTTCTGACTACGACCGTGAAAA 1091

TCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAA

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; FEATURE:

; NAME/KEY: misc_feature

; LOCATION: (1)...(461)

; OTHER INFORMATION: n = A,T,C or G

US-09-404-879A-27
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; TYPE: DNA
; ORGANISM: Escherichia coli
US-09-472-971-7
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APPLICANT: SOGO, KAZUYO
APPLICANT: YANAGI, Hideki
APPLICANT: YURA, Takashi
TITLE OF INVENTION: TRIGGER FACTOR EXPRESSION PLASMIDS
FILE REFERENCE: 1422-409P
                                                                                                                                                                                                                                                                                   SOFTWARE:
SEQ ID NO 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sequence 27, Application US/09404879A
Patent No. 6468546
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                                                                                        Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                APPLICANT: Algate, Paul A.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER FILE REFERENCE: 210121.462C2

CURRENT APPLICATION NUMBER: US/09/404,879A

CURRENT FILING DATE: 1999-09-24

NUMBER OF SEQ ID NOS: 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/472,971
CURRENT FILING DATE: 1999-12-28
EARLIER APPLICATION NUMBER: JP10-372965
EARLIER FILING DATE: 1998-12-28
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SOFTWARE: Patentin Ver
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                                                                                                                                                                                                                                                LENGTH: 46
                                                                                                                                                                                                                             ORGANISM: Homo sapien
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 209
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                                                                                     Local Similarity
                               1 AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 60
AGCAAGGACAATGACTGCTTGAATTGAGGCCTTGAGGAATGAAGCTTTGAAGGAAAAGAA 150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TCCGTCAGCAGATTGAAGAAGCAACTTCTGACTACGACCGTGAAAA 147:
                                                                                                                                                                                                                                                                  461
                                                                                                                                                                                                                                                                                                     FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                              King, Gordon E.
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                                                                      Conservative
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                                                                                     38.7%;
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Pred. No. 9.1;
0; Mismatches
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                                                                                   Score 23.2;
Pred. No. 7.
                                                                    Mismatches
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RESULT 7 US-09-404-879A-74

CORRESPONDENCE ADDRESS:

NUMBER OF SEQUENCES:

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US-09-404-879A-391
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Mitcham, Jennifer L.
APPLICANT: King, Gordon E.
APPLICANT: Algate, Paul A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.462C2
                                                                                                                                                                                    Sequence 139,
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                                                                                                                                                                  Patent No.
                                                                                                                                                                                                                                                                                                                                                           Matches
                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQ ID NO 391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 391, Application US/09404879A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/404,879A
CURRENT FILING DATE: 1999-09-24
NUMBER OF SEQ ID NOS: 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TITLE OF INVENTION: DIAGNOSIS OF OVARIAN CANCER FILE REFERENCE: 210121.462C2
                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 393
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 1999-C
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APPLICANT: Algate, Paul A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 1567
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapien
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 2627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1356 AGCAAGGACAATGACTGCTTGAATTGAGGCCTTGAGGAATGAAGCTTTGAAGGAAAAGAA 1415
                                                      APPLICANT:
                                                                                                                                                                                                                                                                                 2399 AGCAAGGACAATGACTGCTTGAATTGAGGCCTTGAGGAATGAAGCTTTGAAGGAAAAGAA 2458
                                                                                                                APPLICANT:
TITLE OF INVENTION: ENGINEERED HUMAN-DERIVED KUNITZ
TITLE OF INVENTION: DOMAINS THAT INHIBIT HUMAN NEUTROPHIL ELASTASE
                                      APPLICANT:
                                                                                           APPLICANT:
                                                                                                                               APPLICANT:
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nes 37; Conserv
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es 37; Conserv
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                                                                                                                                                                                    Application
                                                                     ROBERTS, Bruce L.
                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                  MARKLAND, Willia KENT, Rachel B.
                                                                                                            LEY, Arthur C. LADNER, Robert C.
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61.7%;
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                                                                                                                                                                                                                                                                                                                                                                        38.7%;
61.7%;
                                                      William
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                                                                                                                                                                                                                                                                                                                                                       Score 23.2; D
Pred. No. 11;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                             DB 4;
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RESULT 10
US-09-138-024-20/c
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Best Local Similarity
                                                                                                                                                                                          GENERAL INFORMATION:
                                                                                                                                                                                                                           Sequence 20,
                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                              APPLICANT: MOOFE, Jeffrey B.
APPLICANT: Wobbe, C. Richard
APPLICANT: Healy, Judith M.
APPLICANT: Donnelly, Caroline E.
TITLE OF INVENTION: REGULATED GENE EXPRESSION IN YEAST
FILE REFERENCE: 0342/10469US1
              CURRENT APPLICATION NUMBER: US/09/138,024A CURRENT FILING DATE: 1998-08-21
                                                                                                                                                      APPLICANT: Bradley, John D. APPLICANT: Thompson, Craig M.
PARLIER APPLICATION NUMBER: 60/056,719
                                                                                                                                                                                                            tent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 202-737-3528
TELEX: 248633
NFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/133,031
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION: TELEPHONE: 202-628-5197
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STREET: 4:
                                                                                                                                                                                                                                                                                                                                   15 GTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAGA 59
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: FILING DATE: 13-003
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: FILING DATE: 02-MA
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                                                                                                                                                                                                                                                                                                               GTGAAAATGTTGAAGAASHAACGSNKGAAGANASCAAYACAANGA 45
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419 Seventh Street, N.W. Suite 300
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02-MAR-1990
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26-JAN-1993
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Pred. No. 11;
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Query Match
Best Local Similarity
Thehes 32; Conserv
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US-08-967-101-5/c
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US-09-404-066-20
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US-09-138-024-20
                                                                                                                                                                Sequence 5, Application US/08967101 Patent No. 5840540
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LENGTH: 7102
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Best Local Similarity
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                                                                                                                                               GENERAL INFORMATION:
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CURRENT APPLICATION NUMBER:
CURRENT FILING DATE: 1999-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Healy, Judith M.
APPLICANT: Donnelly, Caroline E.
TITLE OF INVENTION: REGULATED GENE EXPRESSION IN YEAST
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Bradley, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SQFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 7102
TYPE: DNA
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE:
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                                                                                              APPLICANT: ST. GEORGE-HY
APPLICANT: ROMMENS, JOHA
APPLICANT: FRASER, PAUL
                                   CORRESPONDENCE ADDRESS:
                                                  NUMBER OF SEQUENCES:
                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                 476 ATTATATGCGAAAATTATAAAAGAAAGCAATGAGAAGAAACATTAAAA 429
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   STREET:
                  ADDRESSEE:
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6365409
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Moore, Jeffrey B.
Wobbe, C. Richard
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Thompson,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/09404066
E: TESTA, HURWITZ & THIBEAULT
High Street Tower - 125 High Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                               Conservative
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                                                                                                                                  GEORGE-HYSLOP, PETER H
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1998-08-21
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66.7%;
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                                               GENETIC SEQUENCES AND PROTEINS RELATED TO ALZHEIMER'S DISEASE
                                                                                                                                                                                                                                                                                                                                             37.3%;
                                                                                                                   JOHANNA M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60/056,719
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                                                                                                                                                                                                                                                                                                                                           Score 22.4;
Pred. No. 28;
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Pred. No. 28;
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RESULT 13
US-08-592-541-5/c
; Sequence 5, Application US/08592541
; Sequence 5, Application US/08592541
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                                 CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: PItcher, Edmund R.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 248-7000
TELEPAX: (617) 248-7100
INFORMATION FOR SEQ ID NO: 5:
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INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 3087 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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Best Local S
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APPLICANT:
APPLICANT:
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APPLICANT: FRASER, PAUL
TITLE OF INVENTION: GENER
TITLE OF INVENTION: TO A
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TELECHMUNICATION INFORMATION:
TELEPHONE: (617) 248-7100
TELEPAX: (617) 248-7100
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SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                    SEQUENCE CHARACTERISTICS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     375 CACTATGGATTAATANTGCCTAAAGCAGAGCTGACCACAACGGTGAGAAGAA 324
                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
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                                                                                                                                                                APPLICATION NUMBER: FILING DATE:
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                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
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                                                                                                                                                                                                                                                                                                            02110
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High Street Tower - 125 High Street
3087 base pairs
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SYSTEM: PC-DOS/MS-DOS
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Pred. No. 27;
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RESULT 15
US-09-127-480-5/c
US-09-127-480-5/c
; Sequence 5, Application US/09127480
; Patent No. 6194153
; GENERAL INFORMATION:
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US-09-124-698-5/c
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US-08-592-541-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 5, Application US/09124698 Patent No. 6117978
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 37.0%;
Best Local Similarity 63.5%;
Matches 33; Conservative
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                                                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                         NAME: Pitcher, Edmund R. TELECOMMUNICATION INFORMATION: TELEPHONE: (617) 248-7000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/124,698
                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 3087 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION: NAME: Pitcher, Edmund R.
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PRIOR APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION:
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                                                                                                                375 CACTATGGATTAATANTGCCTAAAGCAGAGCTGACCACAACGGTGAGAAGAA 324
                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY:
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High Street Tower - 125 High Street
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(617) 248-7100
--- TD NO: 5:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ST. GEORGE-HYSLOP, PETER H ROMMENS, JOHANNA M
                                                                                                                                                                               Conservative
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63.5%;
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Pred. No. 27;
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Pred. No. 27;
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CLASSIFICATION:

PRIOR APPLICATION NUMBER: US/08/592,541

PRIOR APPLICATION NUMBER: US/08/592,541

PILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: PItcher, Edmund R.

TELECOMMUNICATION INFORMATION:

TELEPHONE: (617) 248-7100

INFORMATION FOR SEQ ID NO: 5:

SEQUENCE CHARACTERISTICS:

LENGTH: 3087 base pairs

TYPE: nucleic acid

STRANDENNESS: single

STRANDENNESS: single

TOPOLOGY: linear

MOLECULE TYPE: CDNA

US-09-127-480-5
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Search completed: May 24, 2003, 15:37:35 Job time: 27.8387 secs
                                                                                                                                               Query Match 37.0%;
Best Local Similarity 63.5%;
Matches 33; Conservative
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARRE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/127,480
FILING DATE:
FILING DATE:
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TITLE OF INVENTION: TO
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                    375 CACTATGGATTAATANTGCCTAAAGCAGAGCTGACCACAACGGTGAGAAGAA 324
                                                                                              9 CACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACGACAATGAAAAGAA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ST. GEORGE-HYSLOP, PETER H
ROMMENS, JOHANNA M
FRASER, PAUL E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENETIC SEQUENCES AND PROTEINS RELATED
TO ALZHEIMER'S DISEASE
183
                                                                                                                                               Score 22.2; DB 4;
Pred. No. 27;
0; Mismatches 19;
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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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Perfect score:
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    seq
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    828747 seqs, 660231138 residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                             /cgn2_6/ptodata/2/pubpna/US08_NEW_PUB.seq:*
/cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*
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                 462
600
301
371
398
472
541
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532
8340
12241
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    US-09-963-803-21
US-09-963-803-20
US-09-963-803-22
US-09-963-803-7
US-09-963-803-7
US-09-963-803-25
US-09-963-803-25
US-09-963-803-25
US-09-963-803-24
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US-09-963-803-3
US-09-963-803-4
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1693.834 Million cell updates/sec
               Sequence 3
Sequence 4
Sequence 2
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Sequence 7
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ee 4, Appli
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23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.2	23.4	23.6	23.6	23.6	23.6	23.6	23.6	23.6	23.8	23.8	24.2	24.2	24.4
38.7	38.7	38.7	38.7	38.7	38.7	38.7	38.7	38.7	38.7	38.7	38.7	38.7	39.0	39.3	39.3	39.3	39.3	39.3	39.3	39.3	39.7	39.7	40.3	40.3	40.7
555	524	486	461	461	442	426	422	391	352	285	285	281	802	30246	4509	4473	2155	400	373	367	518	272	2259	2259	513509
10	10	10	10	9	10	10	10	10	10	10	9	10	10	7	10	10	10	7	7	10	10	10	10	10	9
US-09-833-790-388	US-09-867-701-2301	US-09-867-701-1482	US-09-884-441-27	US-09-907-969-27	US-09-867-701-2832	US-09-867-701-4161	US-09-867-701-4906	US-09-867-701-3918	US-09-867-701-6377	US-09-777-564-1338	US-10-015-219-1338	US-09-850-178-4	US-09-867-550-1365	US-08-781-986A-56	US-09-815-242-8064	US-09-815-242-4513	US-09-960-428-13	US-08-781-986A-3782	US-08-781-986A-3139	US-09-764-877-65	US-09-864-761-13307	US-09-850-178-31	US-09-815-242-9563	US-09-815-242-9323	US-09-754-853A-4
Sequence 388, App	2301,	Sequence 1482, Ap	Sequence 27, Appl	Sequence 27, Appl	Sequence 2832, Ap			Sequence 3918, Ap	-	Sequence 1338, Ap	Sequence 1338, Ap	Sequence 4, Appli	Sequence 1365, Ap	Sequence 56, Appl	Sequence 8064, Ap	Sequence 4513, Ap	Sequence 13, Appl	Sequence 3782, Ap	Sequence 3139, Ap	Sequence 65, Appl	Sequence 13307, A	Sequence 31, Appl	Sequence 9563, Ap	Sequence 9323, Ap	Sequence 4, Appli

ALIGNMENTS

US-09-963-803-9

Sequence 9, Application US/09963803 Publication No. US20030028922A1 GENERAL INFORMATION:

APPLICANT: Meristem Therapuetics

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PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
LENGTH: 60
TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                               RESULT 2
US-09-963-803-3
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                                                                                                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Directional desoxynucleotide building ; OTHER INFORMATION: S2 US-09-963-803-9
                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches 60
Sequence 3, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello TITLE OF INVENTION: Virus and cassava vein mosaic virus FILE REFERENCE: 18433042 CURRENT APPLICATION NUMBER: US/09/963,803 CURRENT FILING DATE: 2001-09-26 PRIOR APPLICATION NUMBER: F 99/03925 PRIOR APPLICATION NUMBER: F 99/03925 PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
                                                                                                                                                                                                                      1 AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 60
                                                                                                                                                                                                    AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA
                                                                                                                                                                                                                                                                                    60;
                                                                                                                                                                                                                                                                                                             Similarity
                                                                                                                                                                                                                                                                                       Conservative
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Pred. No.
                                                                                                                                                                                                                                                                                       Mismatches
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TITLE OF INVENTION: virus and cassava vein mosaic virus FILE REFERENCE: 184332042

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RESULT 4
US-09-963-803-21
Sequence 21, Application US/09963803
; Sequence No. US20030028922A1
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                                                                                                                                                                                                                                                            ; NAME/KEY: promoter
; LOCATION: (1)..(348)
; OTHER INFORMATION:
US-09-963-803-4
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US-09-963-803-4
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 4, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: Virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05 NUMBER OF SEQ ID NOS: 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: promoter MPr1117
                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: promoter LOCATION: (1)..(317) OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
OTHER INFORMATION: Promoter MPr1116
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                      FEATURE
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                                                                                                                                                                                                                                                                                                                                                                                                                              ENGTH:
                                                                                                                107
                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             73
                                                                                                                                1 AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 60
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                                                                                                           AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AGCCATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAAGAA 132
                                                                                                                                                                                                                                                                                                                                                                                                                                  348
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                                                                                                                                                                                      Conservative
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Pred. No. 1.2e-12;
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RESULT 6

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RESULT 5
US-09-963-803-19
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; LOCATION: (1)..(393)
; OTHER INFORMATION:
US-09-963-803-19
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; LOCATION: (1)..(392)
; OTHER INFORMATION:
US-09-963-803-21
                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION NUMBER: PCT 1800/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
SEQ ID NO 19
LENGTH: 393
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 19, Application US/09963803
Publication No. US20030028922A1
GENERAL INFORMATION:
APPLICANT: Meristem Therapuetics
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APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                           Matches
                                                                                           Query Match
Best Local Similarity
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: ER 99/03925
PRIOR FILING DATE: 1999-03-29
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CURRENT FILLING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
PRIOR FILING DATE: 2000-10-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: virus and cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                  FEATURE:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:
OTHER INFORMATION: promoter MPrll64
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 75
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1 AGCCATGACACTCTGTGCGAATATTGAAGAGCGTAAGCACTGACGACAACAATGAAAAGAA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          60;
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                                                                           Conservative
                                                                                                                                                                                                                                                  promoter MPr1162
                                                                                           100.0%;
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Pred. No. 1.3e-1;
                                                                                           Score 60; DB 9;
Pred. No. 1.3e-12;
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                                                                           Mismatches
                                                                                                             Length 393;
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                                                                             Indels
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                                                                         Gaps
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; OTHER INFORMATION: US-09-963-803-20
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; LOCATION: (1)..(600)
; OTHER INFORMATION:
US-09-963-803-22
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Publication No. US20030028922A1
GENERAL INFORMATION:
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                                                                   Matches
                                                                                                  Query Match
                                                                                                                                                                                                                                                                                   SEQ ID NO 22
LENGTH: 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -09-963-803-22
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION NUMBER: PCT IB00/00370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR FILING DATE: 2000-10-05
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                                                                                                                                                                                                                                                                  TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
NAME/KEY: promoter
LOCATION: (1)..(46
                                                                                                                                                                                                FEATURE:
                                                                                                                                                                                                                OTHER INFORMATION: promoter MPr1165
                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                   FEATURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75
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Local Similarity
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                                                                 Conservative
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                                                                             100.0%;
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                                                                               Score 60;
Pred. No.
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Pred. No.
                                                                 Mismatches
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                                                               1.4e-12;
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                                                               Indels
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; NAME/KEY: promoter
; LOCATION: (1)..(371)
; OTHER INFORMATION:
US-09-963-803-5
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APPLICANT: Meristem Therapuetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
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APPLICANT: Meristem Therapuetics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 5, Application US/09963803 Publication No. US20030028922A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQ ID NO 7
LENGTH: 301
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                                                                                                                                                                                                                                                            SEQ ID NO 5
LENGTH: 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
                Query Match
Best Local
                                                                                                                                                                                                                                                                                                CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
SOFTMARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 184332042
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello TITLE OF INVENTION: virus and cassava vein mosaic virus
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SOFTWARE: PatentIn version 3.1
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PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT 1800/00370
PRIOR FILING DATE: 2000-10-05
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                                                                                                                                                           FEATURE:
OTHER INFORMATION: promoter MPr1146
FEATURE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial Sequence
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  56;
                  Similarity
  Conservative
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                  92.3%;
98.2%;
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                Score 55.4; DB
Pred. No. 6e-11;
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                                     DB 9;
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                                     Length 371;
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Gaps
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US-09-963-803-25
; Sequence 25, Application US/09963803
; Publication No. US20030028922A1
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; LOCATION: (1)..(398)
; OTHER INFORMATION:
US-09-963-803-6
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Query Match
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Best Local Similarity
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TITLE OF INVENTION: chimeric expression promoters originating
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR APPLICATION NUMBER: PCT IB00/00370
NUMBER OF SEQ ID NOS: 39
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TITLE OF INVENTION: Chimeric expression promoters originating
TITLE OF INVENTION: VITUS and cassava vein mosaic virus
FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
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                                                    NAME/KEY: promoter LOCATION: (1)..(472) OTHER INFORMATION:
                                                                                                                                  FEATURE:
OTHER INFORMATION: promoter MPrll69
                                                                                                                                                                            LENGTH: 472
TYPE: DNA
ORGANISM: Artificial Sequence
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TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                         FEATURE:
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92.3%;
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Pred. No. 6.1e-11;
Score 55.4;
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BB
9,
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Length 472;
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                                                            FEATURE
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RESULT 12
US-09-963-803-24
; Sequence 24, Application US/09963803
; Publication No. US20030028922A1
; GENERAL INFORMATION:
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US-09-963-803-23
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Publication No. US20030028922A1
GENERAL INFORMATION:
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SEQ ID NO 23
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                                                                                                                                                                                                                                                                                                                    FILE REFERENCE: 184332042
CURRENT APPLICATION NUMBER: US/09/963,803
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Meristem Therapuetics
TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: VITUS and Cassava vein mosaic virus
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CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: FR 99/03925
PRIOR FILING DATE: 1999-03-29
PRIOR APPLICATION NUMBER: PCT IB00/00370
PRIOR FILING DATE: 2000-10-05
NUMBER OF SEQ ID NOS: 39
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yello
TITLE OF INVENTION: Virus and cassava vein mosaic virus
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                                                                                                                         LENGTH: 604
TYPE: DNA
ORGANISM: Artificial Sequence
NAME/KEY: promoter LOCATION: (1)..(604) OTHER INFORMATION:
                                                                                      FEATURE: OTHER INFORMATION: promoter MPr1167
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TYPE: DNA
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OTHER INFORMATION:
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Pred. No. 6.7e-11;
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); Mismatches 1;
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US-09-963-803-23

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RESULT 15
US-09-765-555-1
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; LOCATION: (1)..(515)
; OTHER INFORMATION:
US-09-963-803-2
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US-09-963-803-2
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Best Local Similarity 97.6
Matches 41; Conservative
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Publication No. US20030037355A1
GENERAL INFORMATION:
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                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 1
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SEQ ID NO 2
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                                                                                                                                                                                                                                                                            APPLICANT: The Scripps Research Institute TITLE OF INVENTION: Methods and compositions to modulate TITLE OF INVENTION: expression in plants FILE REFERICE: 27801-20014.40 (CURRENT APPLICATION NUMBER: US/09/765,555 (CURRENT FILING DATE: 2002-05-24 (PRIOR APPLICATION NUMBER: US 09/620,897 (PRIOR APPLICATION NUMBER: US 09/620,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION NUMBER: PCT IB00/00370 PRIOR FILING DATE: 2000-10-05
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CURRENT FILING DATE: 2001-09-26
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TITLE OF INVENTION: Chimeric expression promoters originating from commelina yellow
TITLE OF INVENTION: virus and cassava vein mosaic virus
FILE REPERENCE: 184332042
                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS:
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                         LENGTH: 532
TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Promoter from the intergenic region of Cassava Vein Mosaic virus OTHER INFORMATION: of 515 bp in length EMBL
OTHER INFORMATION: Promoter CSVMV
                                         ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE
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mes 56; Conserv
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97.6%;
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Pred. No. 6.9e-11;
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Pred. No. 2.1e-05;
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US-09-765-555-1

Query Match 67.3%; Score 40.4; DB 9; Length 532; Best Local Similarity 97.6%; Pred. No. 2.1e-05; Matches 41; Conservative 0; Mismatches 1; Indels 0;

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Search completed: May 24, 2003, 15:35:59
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AZ389667 1M0150L03
AQ633949 RPCI-11-4
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ALIGNMENTS

REFERENCE AUTHORS RESULT 1 AU230643/c LOCUS SOURCE ORGANISM COMMENT ACCESSION DEFINITION KEYWORDS VERSION TITLE JOURNAL Tel: 81-298-36-4359
Fax: 81-298-36-9060
Email: msekl@rtc.riken.go.jp
An Arabidopsis full-length cDNA library was constructed essentially
as reported previously (Seki et al., 1998).cDNA cleaved with BamHI Seki, M., Narusaka, M., Ishida, J., Kamiya, A., Satou, M., Nakajima, M., Akiyama, K., Enju, A., Oono, Y., Sakurai, T., Carninci, P., Kawai, J., Itoh, M., Ishii, Y., Arakawa, T., Shibata, K., Shinagawa, A., Muramatsu, M., Hayashizaki, Y. and Shinozaki, K.
Large scale analysis of Arabidopsis full-length cDNA Unpublished (2002)
Contact: Motoaki Seki
Plant Functional Genomics Research Group
RIKEN Genomic Sciences Center
3-1-1 Koyadai, Tsukuba, Ibaraki 305-0074, Japan AU230643 RAFL19 Arabidopsis thaliana cDNA clone RAFL19-72-A12 3', Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis. mRNA sequence. AU230643 Arabidopsis thaliana thale cress AU230643.1 GI:19799353 (bases 1 to 385) EST 25-APR-2002

AW605446

PM4-DT006

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JOURNAL COMMENT
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                                                            ACTGAGTGAAAAAGTGCAAAATTTAAGCATTAACGACAACAATGAAAA 295
                                                                                                                                                                                                                                                                                                                                                                                                                                              Department of Genetics
Washington University Medical:
Box 8232, 4566 Scott Ave., St.
Tel: 314 362 2735
Fax: 314 362 7855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 (bases 1 to 424)
Cliften, P.F., Hillier, L.W., Fulton, L., Graves, T., Miner, T., Gish, Warerston, R.H. and Johnston, M.
Surveying Saccharomyces genomes to identify functional elements by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Saccharomyces mikatae
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
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AZ924724

AZ926.id55c22.sl Saccharomyces mikatae IFO 1815 Saccharomyces mikatae genomic clone 4906.id55c22.sl, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  and XhoI was ligated to modified Lambda FLC-1 vector (Carninci et al., submitted for publication) digested with BamHI and SalI. This clone is in a modified pBluescript vector. Please visit our web site (http://www.gsc.riken.go.jp/e/plant/index _e.html) for further
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Comparative DNA sequence Unpublished (2001)
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                                                                                                                                                                                                                                                                                                                                                                                                          Email: mj@genetics.wustl.edu
Class: random plasmid subclone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Contact: Johnston M
                                                                                                                                                          Similarity
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                                                                                                                                                                                                                            /db_xref="taxon:114525"
/clone="4906.1055c22.sl"
/clone="10"-"Saccharomyces mikatae
/note="Random genomic sequence"
123 c 80 g 132 t
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/strain="IFO 1815"
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93 c 57 g 114.t
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/db_xref="taxon:3702"
/clone="RaFL19-72-112"
/clone_11b="RaFL19"
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/lab_host="DH10B"
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                                             Mus musculus
Mus musculus
Eukaryota; Metazoa; Chordata;
Eukaryota; Metazoa; Chordata;
                                                                                                                                                                           AZ389667 431 bp DNA 1:
1M0150L03R Mouse 10kb plasmid UUGC1M library
clone UUGC1M0150L03 R, DNA sequence.
Dunn, D., Aoyagi, A., Ba:
Islam, H., Longacre, S.,
                                  Mammalia; Eutheria;
1 (bases 1 to 431)
                                                                                                                                            AZ389667
AZ389667.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Clones are derived from the mouse BAC library RPCI-24. For BAC library availability, please contact Pieter de Jong (pdejong@mail.cho.org). Clones may be purchased from BACPAC Resources (http://www.chorl.org/bacpac/orderingframe.htm). BAC page: http://www.tigr.org/tdb/bac_ends/mouse/bac_end_intro.html
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9712 Medical Center Dr.,
Tel: 301 838 0200
Fax: 301 838 0208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Department of Eukaryotic Genomics
The Institute for Genomic Research
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Tsegaye, G., Geer, K., Krol, M., Shvartsbeyn, A.,
Russell, D., de Jong, P. and Fraser, C.M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mus musculus
Eukaryota; Metazoa;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalia; Eutheria;
                                                                                                         house mouse
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                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /cell_type="Spleen/Brain"
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector: pTARBAC1; Site_1: BamH1; Site_2: BamH1;
/note="Vector pTARBAC1; Site_1: BamH1; Site_2: BamH1;
//note="Vector of the pTARBAC1 cloning vector at the lamH1 sites using Mbol partially digested male C57BL/6J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /strain="C57BL/6J"
/db_xref="taxon:10090"
/clone="RPCI-24-367F11"
/clone_1lb="RPCI-24"
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1. .667
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Rodentia;
Barber, M., Bea
S., Mahmoud, M.,
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Mus musculus
                                                    Craniata; Vertebrata; Euteleostomi; Sciurognathi; Muridae; Murinae; Mus
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              Beacorn, T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 186
 corn,T., Duval,B., Hamil Meenen,E., Pedersen,T.,
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; Murinae; Mus
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e RPCI-24-367F11
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RESULT 5
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                Eukaryota; Metazoa; Cl
Mammalia; Eutheria; P.
1 (bases 1 to 575)
Zhao,S., Adams,M.D.,
Zhao,S.,
                                                                                                                                                                                                   RPCI-11-478I1.TV RPCI-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    High quality sequence stop: 431
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   University of Utah University of Utah
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                                                                                        Homo sapiens
                                                                                                            numan
                                                                                                                                                AQ633949.1 GI:5096584
                                                                                                                                                                                DNA sequence.
                                                                                                                                                                                                                    AQ633949
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mouse whole genome scaffolding with paired end reads from 10kb
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39; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pMP42 (gi14732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheared, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ddunn@genetics.utah.edu
Length: 10000 Std Erro
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/strain="C57BL/6J"
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/clone="UUGC1M0150L03"
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                                                  Metazoa; Chordata; Craniata; Vertebrata; Eutheria; Primates; Catarrhini; Hominidae;
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                  Nierman, W., Malek, J.,
                                                                                                                                                                                                                                                                                                                                                                               Score 26.4; DB 17;
Pred. No. 3.3e+02;
0; Mismatches 21;
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"Omo sapiens"
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Other_GSSs: RPCI-11-478I1.TJ
CONTACT: Shaying Zhao, William Nier
Department of Eukaryotic Genomics
The Institute for Genomic Research
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: hbe@tigr.org

Clones are derived from the human BAC library RPCI-11. For BA

Clones availability, please contact Pieter de Jong

library availability, please contact Pieter de Jong

(pieter@dejong.med.buffalo.edu). Clones may be purchased from

BACPAC Resources (http://bacpac.med.buffalo.edu/ordering) or

Research Genet cs (info@resgen.com). BAC end search page:
                                                                                                                          Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
Project. This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=PM4&t2=PM4-DT0067-010200-002-b05&t3=2000-02-01&t4=1)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Unpublished (1999)
Contact: Simpson A.J.G.
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High quality sequence stop: 118.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                        Tel: +55-11-2704922
Fax: +55-11-2707001
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Fax: 301 838 0208
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/note="Vector: pBACe3.5; Site_1:
/note="Vector: pBACe3.5; Site_1:
/note="Vector: pBACe3.5; Site_1:
/note="Vector: pBACe3.5; Site_1: pBACe3.5; Site_1:
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/cell_type=
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/clone="RPCI-11-478I1"
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/db_xref="GDB:7683360"
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69.2%;
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Pred. No. 3
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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Tel: 801 585 5606
Fax: 801 585 7177
Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Error: C
Plate: 0108 row: G column: 23
Seq primer: CACACAGGAAAACAGCTATGACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  l Similarity
37; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      University of Utah Genome Center University of Utah
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                                                                                                                                                                                                                                                                                                                                                                                                               High quality sequence stop: 658
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        plasmid inserts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and Wright, D., Weiss, R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mouse whole genome scaffolding with paired end reads from 10kb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Dunn, D., Aoyagi, A., Barber, M., Beacorn, T., Duval, B., Hamil, C.,
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M., Rose,M., Rose,R., Stokes,R., Tingey,A., von Niederhausern,A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         lass: plasmid ends
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a 21 c 61 g 56 t 4 others
(http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through a 0.005 inch orifice at constant velocity. The sheared DNA was blunt end-repaired with T4 DNA polymerase and T4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Biomedical Polymers Research Bldg.,
                                                                                                     musculus C57BL/6J (male) was obtained Laboratory Mouse DNA Resource
                                                                                                                                                  /note="Vector: PWD42nv; Purified genomic DNA from M.
                                                                                                                                                                                                                                                          /db_xref="taxon:10090"
/clone="UUGC1M0108G23"
                                                                                                                                                                                                                                                                                                            /organism="Mus musculus"
/strain="C57BL/6J"
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/db_xref="taxon:9606"
                                                                                                                                                                            'lab_host="E. Coli strain XL10-Gold, T1-resistant,
                                                                                                                                                                                                                                  'clone_lib="Mouse 10kb plasmid UUGCIM library"
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Pred. No. 3.4e+02;
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Query Match Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    110ear EST 06-DEC-1999 up05e09, yl NCI_CGAP_Lu30 Mus musculus cDNA clone IMAGE:2651176 5' similar to SW:ERP5_HUMAN Q15084 PROBABLE PROTEIN DISULFIDE ISOMERASE P5 PRECURSOR ;, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA Library Preparation: Life Technologies, Inc. CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL) DNA Sequencing by: Washington University Genome Sequencin Clone distribution: NCI-CGAP clone distribution informatifound through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Gilbert Smith, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                  High quality sequence stop:
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                        Seq primer: -40RP from Gibco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCI-CGAP http://www.ncbi.nlm.nih.gov/ncicgap.
National Cancer Institute, Cancer Genome Anatomy Project (CGAP),
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mammalia; Eutheria;
1 (bases 1 to 540)
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus
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                        /note="Organ: lung; Vector: pCMV-SPORT6; Site_1: NotI; Site_2: SalI; transgenic model WNT-1, expression driven by MMTV-ITR enhancer; Cloned unidirectionally. Primer: Oligo dr. Library constructed by Life Technologies. Investigator providing samples: Gilbert Smith, NIH" 125 c 142 g 113 t 11 others
                                                                                                                                                                                                                                                                                  /organism="Mus musculus"
/strain="CZECH II"
/db_xref="taxon:10090"
                                                                                                                                                                           /clone="IMAGE:2651176"
/clone_11b="NCI_CGAP_Lu30"
/tlssue_type="tumor, metastatic to mammary"
/lab_host="DH10B"
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Pred. No. 4.1e+02;
0; Mismatches 8;
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                                                                                                                                                                                                                                                                                                          CATGACACTCTGTGCGAATATTGAAGACGTAAGCACTGACGACAACAATGAAAA
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Homo sapiens cDNA ADB clones
Unpublished (2000)
Contact: Zeguang Han
Chinese National Human Genome Center at Shanning Road The
                                 AL322301.1 GI:9555185
GSS; genome survey sequence.
Tetraodon nigroviridis.
Tetraodon nigroviridis
Tetraodon; Ohordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei
Acanthomorpha; Acanthomorpha; Acanthomorpha; Perromorpha; Neopterygii; Percomorpha; Tetraodontiformes;
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037C06 of library A fro
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1 (bases 1 to 598)
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             (bases 1 to 1085)
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lone is available at CHGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /tissue_type="Adrenal gland"
/dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /clone_lib="ADB"
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/db_xref="taxon:9606"
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                            Tetraodon.
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 Jaillon, O.,
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Pred. No. 5.3e
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Dasilva,C.,
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.es 18;
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Bouneau, L.,
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 Fisher, C.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   scale clone-end sequencing project of the Tetraodon genome. For more information, please take a look at http://www.genoscope.cns.fr/Tetraodon.
   Tel: +44 131 650 6760
Fax: +44 131 670 5450
Email: mark.blaxter@ed.ac.uk
The library was prepared by Michelle Lizotte-Waniewski for Alan
Scott, Johns Hopkins University Medical School, Baltimore.
Sequencing was performed by the Pathogen Sequencing Unit, Sanger
Sequencing was performed by the Pathogen Sequencing Unit, The
                                                                                                                                                      Unpublished (2000)
Contact: Blaxter ML
Institute of Cell, Animal and
University of Edinburgh
                                                                                                                                                                                                                                                                                                                  Ascaris suum
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Crollius, H.R., Jaillon, O.,
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Saurin, W.
                                                                                                                                                                                                                                    Blaxter, M.L., Parkinson, J., N., Quayle, M. and Barrell,
                                                                                                                                                                                                                                                                                                                                                                                                              As_tgz_67F12_SKPL Ascaris
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Direct Submission
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                                                                                                                                                                                                                                                                                                                                    pig roundworm.
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33; Conservative
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70.2%;
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a PolyA tail (trimmed)
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Pred. No. 5.8e+02;
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Billault,A., Quetier,F., Saurin,
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LOCUS
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                                                                                                                                                                                                                                                                                                                                                                                                                      Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BF681368
602156557F1 NIH_MGC_83
                                                                                                                                                                                                                                                                                                                               http://image.llnl.gov
Plate: LLCM1151 row: b column: 16
High quality sequence stop: 593.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EST
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BF681368
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BF681368.1 GI:11955263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Plate: 67 row:
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FORWARD: T3
                                                                                                                                                                                                                                                                                                                                                                                                found through the I.M.A.G.E. Consortium/LLNL at:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    94
/Glone_lib="NIH_MGC_83"
/lab_host="hH10B (Tl phage-resistant)"
/note="Organ: prostate; Vector: pDNR-LIB (Clontech);
/note="Organ: prostate; Vector: pDNR-LIB (Clontech);
Site_1: SfiI (ggccgcctcggcc); Site_2: SfiI (ggccattatggcc);
S' and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor
sequence: 5'-ATTCTAGAGGCCGAGGCGCGCAATG-dT[30]BN-3'
(where B = A, C, or G and N = A, C, G, or T). Average
insert size 1.4 kb (range 0.5-4.0 kb). 14/15 colonies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note="Vector: Lambda Zap II; Site_1: EcoRI; Site_2: XhoI; Library was made from dissected teštis germinal zone from adult male Ascaris suum collected from abbatoirs.

Constructed by Michelle Lizotte-Waniewski for Alan Scott, Johns Hopkins University Medical School, Baltimore, MD."

a. 97 c. 63 g. 202 t.
                                                                                                                                                                                                              /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="IMAGE:4297263"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism-"Ascaris suum"
/db_xref="taxon:6253"
/clone-"As_tgz_67F12"
/clone_11b-"Ascaris suum adult male testis germinal zone
from Alan Scott"
                                                                                                                                                                                                                                                                                                          Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /sex="Male"
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Pred. No. 5.9e+02;
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RESULT 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Clemson University
100 Jordan Hall, Cl
Tel: 864 656 7288
Fax: 864 656 4293
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  for barley genomics: Morex pre-anthesis spike cDNA library Unpublished (2001)
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Y., Henry,D., Palmer,M., Rambo,T., Simmons,J., Choi,D.W., Fenton
,R.D., Close,S.J., Oates,R. and Main,D.
Development of a genetically and physically anchored EST resource
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae; Pooldeae; Triticeae; Hordeum.

1. (bases 1 to 930)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seq primer: AATTAACCCTCACTAAAGGG High quality sequence stop: 590.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Wing RA
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Total hq bases = 465
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/note="Vector: lambdaZAP; Site_1: EcoR1; Site_2: Xho1; Plants were grown in the greenhouse at the University of California, Riverside (Fenton, SJ Close, TJ Close). Whole spike with awns trimmed were collected at white, green and yellow anther stages (Fenton). Total RNA was prepared from each pool, equal quantities of all three RNA pools were combined, poly(A) RNA was purified from the mixture, one primary unamplified cDNA library was made, and 1 million pfu were in vivo excised to give pBluescript SK(-) cDNA phagemids. These steps were performed in the TJ Close lab (Choi) at the University of California, Riverside. Phagemids were plated and picked at the Clemson University Genomics Institute (CUGI) (Begum, Palmer, Frisch, Atkins and Wing) Plasmid DNA preparations, DNA sequencing and sequence analysis were performed at CUGI (Wing, Yu, Frisch, Henry, Simmons, Oates, Rambo, Main). The sequence has
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                                                                                                                                                                                                                                                                                                                                                                                                                                         /clone_lib="Hordeum vulgare pre-anthesis spike EST library HVCDNA0008 (white to yellow anther)" /tissue_type="pre-anthesis spike" /lab_host="SOLR"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone-"HVSMEg0005I11f"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /db_xref="taxon:4513"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /organism="Hordeum vulgare"
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66.18;
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and was constructed by Clontech
Alto, CA)."
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vector

sequence and

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RESULT 14
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LOCUS
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BASE COUNT
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Song F M. and Goodman, R.M.
Molecular cloning of differentially expressed cDNAs associated with
systemic acquired resistance in rice
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Contact: Song, F.M.; Goodman, R.M. Department of Plant Pathology University of Wisconsin-Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BI118726 284 bp mRNA linear EST 26-JUN-2001 EST112 Differentially expressed cDNA libraries of BTH-treated/blast fungus-infected rice leaf tissues Oryza sativa cDNA clone BNHN-n7,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Benzothiadiazole-inducible
Insert Length: 284 Std Error: 0.00
Seg primer: M13 Forward and reverse.
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Fax: 608 262 8643
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oryza sativa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               fmsong@plantpath.wisc.edu; rgoodman@facstaff.wisc.edu
                                                                                                                                                                                                                                                   /Cline="Yuanfengzao (a susceptible variety to rice blast disease)"
/db_xref="taxon:4530"
/clone="BNHN-n7"
/clone=lib="Differentially expressed cDNA libraries of BTH-treated/blast fungus-infected rice leaf tissues"
/tissue_type="Seedling leaves"
/fissue_type="Seedling leaves"
/dev_stage="Three-week-old rice seedlings pretreated with BTH and/or inoculated with Magnaporthe grisea"
/note="Vector: pGEM-T Easy; Rice seedlings were pretreated by foliar spraying with 0.3 mM_benzothiadazole (BTH) and the incomplete were pretreated with seedlings were pretreated by foliar spraying with 0.3 mM_benzothiadazole (BTH) and the incomplete were pretreated with the seedlings were pretreated by foliar spraying with 0.3 mM_benzothiadazole (BTH) and the seedlings were pretreated with the seedling
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                                                             then inoculated with Magnaporthe grisea three days after BTH treatment. Leaf samples were collected twenty-four hour after inoculation and mRNAs were used for construction of the differentially expressed cDNA libraires by suppression subtractive hybridization."
                        libraires by suppression subtractive 77 c 65 g 62 t
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Query Match

42.38;

Score 25.4;

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Search completed: May 24, 2003, 15:33:20 Job time: 793.387 secs

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JOURNAL
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Local
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Ouryza sativa
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
                                                                                                                                                                                                                                                                                                                                                    Contact: Dong HT
Laboratory of Functional Genetics
Bio-technology Institute of Zhejiang University
Kaixuan Road 268*, Hangzhou, Zhejiang, P.R.China
Tel: 0086-571-86892051
                                                                                                                                                                                                                                                                                                                                                                                                                              l (bases 1 to 489)
Dong,H.T., Li,D.B., Zhuang,X.F., Dai,C.G.,
H.F., Jiang,Y.X., Yu,F.C., Gao,Q.K. and Lo
A Gene Expression Screen in Oryza sativa
Unpublished (2001)
                                                                                                                                                                                                                                                                                                          Email: htdong@zjuem.zju.edu.cn
Seq primer: M13 forward primer
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                                                                         Similarity
                                                                                                                                        134
                                                            Conservative
                                                                                                                                                                                                  M.grisea"
                                                                                                                                                                                                                                /organism="Oryza sativa"
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                                                                                                                                     /tissue_type="leaf"
/dev_stage="Mature stage"
/note="Vector: pSport2"
108 c 131 g 116
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Location/Qualifiers
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                                                           Score 25.4; DB 13;
Pred. No. 6.8e+02;
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